



SEQUENCE LISTING

<110> RASTELLI, LUCA
GERRITSEN, MARY

<120> ANGIOGENESIS ASSOCIATED PROTEINS AND NUCLEIC ACIDS
ENCODING THE SAME

<130> 10716/35

<140> 09/815,379

<141> 2001-03-22

<150> 60/191,134

<151> 2000-03-22

<160> 17

<170> PatentIn Ver. 2.1

<210> 1

<211> 2577

<212> DNA

<213> Homo sapiens

<400> 1

ctggcctaga	tactacaact	gaactttttt	tcttttttagt	tactccacag	gatccgctga	60
acataggatg	ttgccacaaa	atctacctcg	tgtatttttc	tctttcactc	atgagctgca	120
caattgcaga	tttgagcaca	atgtctgcag	actgtgttga	aaaactctga	agaacctaat	180
taacacagga	tgacctagga	gtgattctaa	gtctgtgtaa	caagatatta	ctcattagt	240
aatgtgtcag	tcttggtact	gaatgctgca	gataacagca	agtaggttct	cctttatttc	300
tgaagtattc	acttgacctt	ccatcagtaa	gacggacttt	tctaactctgt	tcctggagat	360
attaatggaa	tacagtcatg	tccactcaag	acgagaggca	gatcaatact	gaatatgctg	420
tgtcattgtt	ggaacagttg	aaactgtttt	atgaacagca	gttgtttact	gacatagtgt	480
taattgttga	gggcactgaa	ttcccttgct	ataagatggg	tcttgcaaca	tgtagctctt	540
atttcagggc	catgtttatg	agtggactaa	gtgaaagcaa	acaaacccat	gtacacctga	600
ggaatgtcga	tgctgccacc	ttacagataa	taataactta	tgcatacacg	ggtaacttgg	660
caatgaatga	cagcactgta	gaacagcttt	atgaaacagc	ttgcttccta	caggtagaag	720
atgtgttaca	acgttgtcga	gaatatttaa	ttaaaaaaat	aaatgcagag	aattgtgtac	780
gattgtttag	ttttgctgat	ctcttcagtt	gtgaggaatt	aaaacagagt	gctaaaagaa	840
tggtggagca	caagttcact	gctgtgtatc	atcaggacgc	gttcatgcag	ctgtcacatg	900
acctactgat	agatattctc	agtagtgaca	atttaaattgt	agaaaaggaa	gaaaccgttc	960
gagaagctgc	tatgtctgtg	ctagagtata	acacagaatc	acgatcccag	tatttgtctt	1020
ctgttcttag	ccaaatcaga	attgatgcac	tttcagaagt	aacacagaga	gcttggtttc	1080
aaggtctgcc	acccaatgat	aagtcagtgg	tggttcaagg	tctgtataag	tccatgcccc	1140
agtttttcaa	accaagactt	gggatgacta	aagaggaaat	gatgattttc	attgaagcat	1200
cttcagaaaa	tccttgtagt	ctttactctt	ctgtctgtta	cagcccccaa	gcagaaaaag	1260
tttacaagtt	atgtagccca	ccagctgatt	tgcataagg	tgggaccgtt	gtaactcctg	1320
ataatgatat	ctacatagca	gggggtcaag	ttcctctgaa	aaacacaaaa	acaaatcaca	1380
gtaaaacaag	caaacttcag	actgccttca	gaactgtgaa	ttgcttttat	tggtttgatg	1440
cacagcaaaa	tacctggttt	ccaaagaccc	caatgctttt	tgtccgcata	aagccatctt	1500
tggtttgctg	tgaaggctat	atctatgcaa	ttggaggaga	tagcgtaggt	ggagaactta	1560
atcggaggac	cgtagaaaga	tacgacactg	agaaagatga	gtggacgatg	gtaagccctt	1620
taccttgtgc	ttggcaatgg	agtgcagcag	ttgtggttca	tgactgcatt	tatgtgatga	1680
cactgaacct	catgtactgt	tattttccaa	ggctctgactc	atgggtagaa	atggccatga	1740
gacagactag	taggtccttt	gcttcagctg	cagcttttgg	tgataaaaatt	ttctatattg	1800
gagggttgca	tattgtacc	aattccggca	taagactccc	ctctggcact	gtagatgggt	1860
cttcagtaac	tgtggaaatt	tatgatgtga	ataaaaaatga	gtggaaaatg	gcagccaaca	1920

```

tccctgctaa gaggtactct gacccctgtg ttagagctgt tgtgatctca aattctctat 1980
gtgtgtttat gcgagaaacc cacttaaagt agcgagctaa atacgtcacc taccaatatg 2040
acctggaact tgaccggtgg tctctgcggc agcatatata tgaacgtgta ctgtgggact 2100
tggggagaga ttttcgatgc actgtgggga aactctatcc atcctgcctt gaagagtctc 2160
catggaaacc accaacttat cttttttcaa cggatgggac agaagagttt gaactggatg 2220
gagaaatggg tgcactacca cctgtatagt ggggaagttc agggagtgca cgcctgagtt 2280
atgtgctttg tcattttctt tgctaaacaa aagaggctat gaaagaacta aatatgagta 2340
cataaaattc tatctttgat aaattttatt tttatgccct acttaatat tgcatacagta 2400
taatataat cagtgagtct tacagaaaga tatgcttcca taatatgaaa tagattattc 2460
aataattgag aaactttatg tgtaatcatg agagtataag aatctggatt atctaacatt 2520
gttagccctg tgtatgtaca gttcaaaaag ttcatttata aaagtagttt cctgttc 2577

```

<210> 2

<211> 623

<212> PRT

<213> Homo sapiens

<400> 2

```

Met Ser Thr Gln Asp Glu Arg Gln Ile Asn Thr Glu Tyr Ala Val Ser
  1                      5                      10                      15

Leu Leu Glu Gln Leu Lys Leu Phe Tyr Glu Gln Gln Leu Phe Thr Asp
          20                      25                      30

Ile Val Leu Ile Val Glu Gly Thr Glu Phe Pro Cys His Lys Met Val
          35                      40                      45

Leu Ala Thr Cys Ser Ser Tyr Phe Arg Ala Met Phe Met Ser Gly Leu
          50                      55                      60

Ser Glu Ser Lys Gln Thr His Val His Leu Arg Asn Val Asp Ala Ala
          65                      70                      75                      80

Thr Leu Gln Ile Ile Ile Thr Tyr Ala Tyr Thr Gly Asn Leu Ala Met
          85                      90                      95

Asn Asp Ser Thr Val Glu Gln Leu Tyr Glu Thr Ala Cys Phe Leu Gln
          100                     105                     110

Val Glu Asp Val Leu Gln Arg Cys Arg Glu Tyr Leu Ile Lys Lys Ile
          115                     120                     125

Asn Ala Glu Asn Cys Val Arg Leu Leu Ser Phe Ala Asp Leu Phe Ser
          130                     135                     140

Cys Glu Glu Leu Lys Gln Ser Ala Lys Arg Met Val Glu His Lys Phe
          145                     150                     155                     160

Thr Ala Val Tyr His Gln Asp Ala Phe Met Gln Leu Ser His Asp Leu
          165                     170                     175

Leu Ile Asp Ile Leu Ser Ser Asp Asn Leu Asn Val Glu Lys Glu Glu
          180                     185                     190

Thr Val Arg Glu Ala Ala Met Leu Trp Leu Glu Tyr Asn Thr Glu Ser
          195                     200                     205

```

Arg Ser Gln Tyr Leu Ser Ser Val Leu Ser Gln Ile Arg Ile Asp Ala
 210 215 220
 Leu Ser Glu Val Thr Gln Arg Ala Trp Phe Gln Gly Leu Pro Pro Asn
 225 230 235 240
 Asp Lys Ser Val Val Val Gln Gly Leu Tyr Lys Ser Met Pro Lys Phe
 245 250 255
 Phe Lys Pro Arg Leu Gly Met Thr Lys Glu Glu Met Met Ile Phe Ile
 260 265 270
 Glu Ala Ser Ser Glu Asn Pro Cys Ser Leu Tyr Ser Ser Val Cys Tyr
 275 280 285
 Ser Pro Gln Ala Glu Lys Val Tyr Lys Leu Cys Ser Pro Pro Ala Asp
 290 295 300
 Leu His Lys Val Gly Thr Val Val Thr Pro Asp Asn Asp Ile Tyr Ile
 305 310 315 320
 Ala Gly Gly Gln Val Pro Leu Lys Asn Thr Lys Thr Asn His Ser Lys
 325 330 335
 Thr Ser Lys Leu Gln Thr Ala Phe Arg Thr Val Asn Cys Phe Tyr Trp
 340 345 350
 Phe Asp Ala Gln Gln Asn Thr Trp Phe Pro Lys Thr Pro Met Leu Phe
 355 360 365
 Val Arg Ile Lys Pro Ser Leu Val Cys Cys Glu Gly Tyr Ile Tyr Ala
 370 375 380
 Ile Gly Gly Asp Ser Val Gly Gly Glu Leu Asn Arg Arg Thr Val Glu
 385 390 395 400
 Arg Tyr Asp Thr Glu Lys Asp Glu Trp Thr Met Val Ser Pro Leu Pro
 405 410 415
 Cys Ala Trp Gln Trp Ser Ala Ala Val Val Val His Asp Cys Ile Tyr
 420 425 430
 Val Met Thr Leu Asn Leu Met Tyr Cys Tyr Phe Pro Arg Ser Asp Ser
 435 440 445
 Trp Val Glu Met Ala Met Arg Gln Thr Ser Arg Ser Phe Ala Ser Ala
 450 455 460
 Ala Ala Phe Gly Asp Lys Ile Phe Tyr Ile Gly Gly Leu His Ile Ala
 465 470 475 480
 Thr Asn Ser Gly Ile Arg Leu Pro Ser Gly Thr Val Asp Gly Ser Ser
 485 490 495
 Val Thr Val Glu Ile Tyr Asp Val Asn Lys Asn Glu Trp Lys Met Ala
 500 505 510

Ala Asn Ile Pro Ala Lys Arg Tyr Ser Asp Pro Cys Val Arg Ala Val
 515 520 525

Val Ile Ser Asn Ser Leu Cys Val Phe Met Arg Glu Thr His Leu Asn
 530 535 540

Glu Arg Ala Lys Tyr Val Thr Tyr Gln Tyr Asp Leu Glu Leu Asp Arg
 545 550 555 560

Trp Ser Leu Arg Gln His Ile Ser Glu Arg Val Leu Trp Asp Leu Gly
 565 570 575

Arg Asp Phe Arg Cys Thr Val Gly Lys Leu Tyr Pro Ser Cys Leu Glu
 580 585 590

Glu Ser Pro Trp Lys Pro Pro Thr Tyr Leu Phe Ser Thr Asp Gly Thr
 595 600 605

Glu Glu Phe Glu Leu Asp Gly Glu Met Val Ala Leu Pro Pro Val
 610 615 620

<210> 3

<211> 1749

<212> DNA

<213> Homo sapiens

<400> 3

caaggggagcg aggggtgtcgt agaggggcaga atgaacaaga agaattagga gggagggtgc 60
 gtgtgcccggg gctaggggct ggaagtcctg gctctagtgt cacctcggaa ggaaaaggca 120
 aacagaggag ggaaggcgct ttaggactgc ctggatccag agcactttcc tcggcctcta 180
 caggcctgtg tcgctatggg ttcccccgcc gccccggagg gagcgctggg ctacgtccgc 240
 gagttcactc gccactcctc cgacgtgctg ggcaacctca acgagctgcg cctgcgcggg 300
 atcctcactg acgtcacgct gctgggtggc gggcaacccc tcagagcaca caaggcagtt 360
 ctcatcgctt gcagtggctt cttctattca attttccggg gccgtgcggg agtcgggggtg 420
 gacgtgctct ctctgcccgg gggtcgccga gcgagaggct tcgcccctct attggacttc 480
 atgtacactt cgcgcctgcg cctctctcca gccactgcac cagcagtcct agcggccgcc 540
 acctatttgc agatggagca cgtggtccag gcactgccacc gcttcatcca ggccagctat 600
 gaacctctgg gcatctccct gcgccccctg gaagcagaac ccccaacacc cccaacggcc 660
 cctccaccag gtagtcccag gcgctccgaa ggacacccag acccacctac tgaatctcga 720
 agctgcagtc aaggcccccc cagtccagcc agccctgacc ccaaggcctg caactggaaa 780
 aagtacaagt acatcgtgct aaactctcag gcctcccaag cagggagcct ggtcgggggag 840
 agaagttctg gtcaaccttg cccccaagcc aggtcctcca gtggagacga ggcctccagc 900
 agcagcagca gcagcagcag cagcagcagt gaagaaggac ccattcctgg tccccagagc 960
 aggtctcttc caactgctgc cactgtgcag ttcaaagtgt gggctccagc cagtaccccc 1020
 tacctcctca catcccaggc tcaagacacc tctggatcac cctctgaacg ggctcgtcca 1080
 ctaccgggga gtgaattttt cagctgccag aactgtgagg ctgtggcagg gtgctcatcg 1140
 gggctggact ccttggttcc tggggacgaa gacaaacctt ataagtgtca gctgtgccgg 1200
 tcttcgttcc gctacaaggg caaccttgcc agtcaccgta cagtgcacac aggggaaaag 1260
 ccttaccact gctcaatctg cggagcccgt tttaaccggc cagcaaacct gaaaacgcac 1320
 agccgcatcc attcgggaga gaagccgtat aagtgtgaga cgtgcggctc gcgctttgta 1380
 caggtacgga gccagcctcc aagtggcttc caaggcaaac ctgcaagagg tgggggtgggc 1440
 caaaaggagg ggttctgttc ctcccagagg caggacttga agtctcctcc ctcccagggtg 1500
 gcacatctgc gggcgacagt gctgatccac accggggaga agccctaccc ttgccctacc 1560
 tgcggaaccc gcttcgcgca cctgcagacc ctcaagagcc acgttcgcat ccacaccgga 1620
 gagaagcctt accactgcga cccctgtggc ctgcatttcc ggcacaagag tcaactggcg 1680
 ctgcattctg gccagaaaca cggagctgct accaacacca aagtgcacta ccacattctc 1740
 gggggggccc 1749

<210> 4
 <211> 518
 <212> PRT
 <213> Homo sapiens

<400> 4

```

Met Gly Ser Pro Ala Ala Pro Glu Gly Ala Leu Gly Tyr Val Arg Glu
  1              5              10              15

Phe Thr Arg His Ser Ser Asp Val Leu Gly Asn Leu Asn Glu Leu Arg
      20              25              30

Leu Arg Gly Ile Leu Thr Asp Val Thr Leu Leu Val Gly Gly Gln Pro
      35              40              45

Leu Arg Ala His Lys Ala Val Leu Ile Ala Cys Ser Gly Phe Phe Tyr
      50              55              60

Ser Ile Phe Arg Gly Arg Ala Gly Val Gly Val Asp Val Leu Ser Leu
      65              70              75              80

Pro Gly Gly Pro Glu Ala Arg Gly Phe Ala Pro Leu Leu Asp Phe Met
      85              90              95

Tyr Thr Ser Arg Leu Arg Leu Ser Pro Ala Thr Ala Pro Ala Val Leu
      100             105             110

Ala Ala Ala Thr Tyr Leu Gln Met Glu His Val Val Gln Ala Cys His
      115             120             125

Arg Phe Ile Gln Ala Ser Tyr Glu Pro Leu Gly Ile Ser Leu Arg Pro
      130             135             140

Leu Glu Ala Glu Pro Pro Thr Pro Pro Thr Ala Pro Pro Pro Gly Ser
      145             150             155             160

Pro Arg Arg Ser Glu Gly His Pro Asp Pro Pro Thr Glu Ser Arg Ser
      165             170             175

Cys Ser Gln Gly Pro Pro Ser Pro Ala Ser Pro Asp Pro Lys Ala Cys
      180             185             190

Asn Trp Lys Lys Tyr Lys Tyr Ile Val Leu Asn Ser Gln Ala Ser Gln
      195             200             205

Ala Gly Ser Leu Val Gly Glu Arg Ser Ser Gly Gln Pro Cys Pro Gln
      210             215             220

Ala Arg Leu Pro Ser Gly Asp Glu Ala Ser Ser Ser Ser Ser Ser Ser
      225             230             235             240

Ser Ser Ser Ser Ser Glu Glu Gly Pro Ile Pro Gly Pro Gln Ser Arg
      245             250             255

Leu Ser Pro Thr Ala Ala Thr Val Gln Phe Lys Cys Gly Ala Pro Ala
      260             265             270

```

Ser Thr Pro Tyr Leu Leu Thr Ser Gln Ala Gln Asp Thr Ser Gly Ser
 275 280 285
 Pro Ser Glu Arg Ala Arg Pro Leu Pro Gly Ser Glu Phe Phe Ser Cys
 290 295 300
 Gln Asn Cys Glu Ala Val Ala Gly Cys Ser Ser Gly Leu Asp Ser Leu
 305 310 315 320
 Val Pro Gly Asp Glu Asp Lys Pro Tyr Lys Cys Gln Leu Cys Arg Ser
 325 330 335
 Ser Phe Arg Tyr Lys Gly Asn Leu Ala Ser His Arg Thr Val His Thr
 340 345 350
 Gly Glu Lys Pro Tyr His Cys Ser Ile Cys Gly Ala Arg Phe Asn Arg
 355 360 365
 Pro Ala Asn Leu Lys Thr His Ser Arg Ile His Ser Gly Glu Lys Pro
 370 375 380
 Tyr Lys Cys Glu Thr Cys Gly Ser Arg Phe Val Gln Val Arg Ser Gln
 385 390 395 400
 Pro Pro Ser Gly Phe Gln Gly Lys Pro Ala Arg Gly Gly Val Gly Gln
 405 410 415
 Lys Gly Gly Phe Cys Ser Ser Gln Arg Gln Asp Leu Lys Ser Pro Pro
 420 425 430
 Ser Gln Val Ala His Leu Arg Ala His Val Leu Ile His Thr Gly Glu
 435 440 445
 Lys Pro Tyr Pro Cys Pro Thr Cys Gly Thr Arg Phe Arg His Leu Gln
 450 455 460
 Thr Leu Lys Ser His Val Arg Ile His Thr Gly Glu Lys Pro Tyr His
 465 470 475 480
 Cys Asp Pro Cys Gly Leu His Phe Arg His Lys Ser Gln Leu Arg Leu
 485 490 495
 His Leu Arg Gln Lys His Gly Ala Ala Thr Asn Thr Lys Val His Tyr
 500 505 510
 His Ile Leu Gly Gly Pro
 515

<210> 5

<211> 3501

<212> DNA

<213> Homo sapiens

<220>

<221> modified_base

<222> (3070)

<223> a, t, c, g, other or unknown

<220>

<221> modified_base

<222> (3087)

<223> a, t, c, g, other or unknown

<400> 5

tctttttcct	cgcgctccttt	gccccggaag	tgctcttaca	acattggctg	ccggcggtgac	60
tttgaccgct	tcccgggtgcg	ttaccggcag	ctgaacccac	ccggcgccac	gggactttga	120
cgcggtgctct	gcgcttgcca	tgagactcct	gggagctgca	gccgtcgcgg	ctctggggcg	180
cgaaggggcc	cccgcctccc	taggctggca	gaggaagcag	gttaattgga	aggcctgccg	240
atggtctttca	tcagggtgga	ttcctaata	aaaaatacga	aatattggaa	tctcagctca	300
cattgattct	gggaaaacta	cattaacaga	acgagtcctt	tactacactg	gcagaattgc	360
aaagatgcat	gaggtgaaag	gtaaagatgg	agttggtgct	gtcatggatt	ccatggaact	420
agagagacaa	agaggaatca	ctattcagtc	agcagccact	ttcaccatgt	ggaaagatgt	480
caatattaac	attatagata	ctcctgggca	tgtggacttc	acaatagaag	tggaaagggc	540
cctgagagtg	ttggatgggtg	cagtccttgt	tctctgtgct	gttggaggggg	tacagtgccca	600
gaccatgact	gtcaatcgtc	agatgaagcg	ctacaacggt	ccgttttctaa	ctttttattaa	660
caaattggac	cgaatgggct	ccaacccagc	cagggccctg	cagcaaatga	ggtctaaact	720
aaatcataat	acagcggtta	tgcagatacc	catgggtttg	gagggtaatt	ttaaagggtat	780
tgtagatctt	attgaggaa	gagccatcta	ttttgatgga	gacttttagtc	agattgttcg	840
atatggtgag	attccagctg	aattaagggc	ggcgggccact	gaccaccggc	aggagctaata	900
tgaatgtgtt	gccaatccag	atgaacagct	tggtgagatg	tttctggaag	aaaaaatccc	960
ctcgattttct	gattttaaagc	tagcaattcg	aagagctact	ctgaaaagat	catttactcc	1020
tgtattttttg	ggaagcgctt	tgaagaacaa	aggagttcag	cctcttttag	atgctgtttt	1080
agaatacctc	ccaaatccat	ctgaagtcca	gaactatgct	attctcaata	aaaaggatga	1140
ctcaaaagag	aaaacccaaa	tcctaataga	ctccagtaga	cacaattccc	acccatttgt	1200
aggcctggct	tttcccctgg	aggtaggtcg	atttggacaa	ttaacttatg	ttcgcagtta	1260
tcagggagag	ctaaagaagg	gtgacaccat	ctataacaca	aggacaagaa	agaaagtacg	1320
gttgcaacgg	ctggctcgca	tgcatgccga	catgatggag	gcaagtacag	aggaagtata	1380
tgccggagac	atctgtgcat	tggttggtcat	tgactgtgct	agtggagaca	cattcacaga	1440
caaagccaac	agcggccttt	ctatggagtc	aattcatggt	cctgatcctg	tcatttcaat	1500
agcaatgaag	ccttctaaca	agaacgatct	ggaaaaat	tcaaaaggta	ttggcagggt	1560
tacaagagaa	gatcccat	ttaaagtata	ctttgacact	gagaacaaag	agacagttat	1620
atctggaatg	ggagaattac	acctggaaat	ctatgctcag	aggctggaaa	gagagtatgg	1680
ctgtccttgt	atcacaggaa	agccaaaagt	tgcccttcga	gagaccatta	ctgcccctgt	1740
cccgtttgac	tttacacata	aaaaacaatc	agggtgtgca	ggccagtatg	gaaaagtaat	1800
agggtgtcctg	gagcctctgg	accagagga	ctacactaaa	ttggaatttt	cagatgaaac	1860
attcggatca	aatattccaa	agcagtttgt	gcctgctgta	gaaaaggggt	ttttagatgc	1920
ctgcgagaag	ggccctcttt	ctggtcacaa	gctctctggg	ctccggtttg	tcctgcaaga	1980
tggagcacac	cacatggttg	attctaata	aatctctttc	atccgagcag	gagaagggtg	2040
tcttaaaca	gccttgga	atgcaacatt	atgtattctt	gaacctatta	tggctgtgga	2100
agttgtagct	ccaaatgaat	ttcagggaca	agtaattgca	ggaattaacc	gacgccatgg	2160
ggtaatcact	gggcaagatg	gagttgagga	ctattttaca	ctgtatgcag	atgtccctct	2220
aaatgatatg	tttgggtatt	ccactgaact	taggtcatgc	acagagggaa	agggagaata	2280
cacaatggag	tatagcaggt	atcagccatg	tttaccatcc	acacaagaag	acgtcattaa	2340
taagtatttg	gaagctacag	gtcaacttcc	tgtaaaaaaa	ggaaaagcca	agaactaact	2400
ttgcttactg	tgagttgact	gactctaatt	gaatctgcgt	ggttttgata	ctttgatgga	2460
ttccagtggg	ataaattcag	gctgctgaaa	caagaaattc	tgagcccagg	aagcgggctc	2520
ttctttcttc	aaaagaagcc	cttcttggtc	atattcagga	gcttctgtta	tattcaaagg	2580
taattctatg	tctatctcaa	ctctattgat	tggttttata	gttcattgaa	aatcctcaaa	2640
taaaatataa	ttattactga	aatatgttta	atattttaagg	ggaaaagaga	ctaatttcag	2700
ttatactttt	aagcttagaa	tgtatgttca	tttccaaatt	ttgtatcata	agagttttca	2760
acatagagaa	aagctgaaaa	aatgcaaaaga	ataaccacat	actttccatc	taccttctct	2820
tggttaacggg	ttgtttatca	tataataatt	tgttttgtca	tattttgctt	cactgtctat	2880
tatctgttta	agtcctataa	ctctattttt	agtttgctga	agacttgaaa	gtgaatcgca	2940
tatatcatga	cacttcttgg	agtgctcatta	atgggcaggc	ttttctgttg	aagagtggat	3000

```

tccgatatgtt cttcatagag agtggttttc agattcttca ttgggatatt aaaatattag 3060
ccaaatttcn ctctgtttta tatatgncag tttatttcag tttgtgggtt ctgcaaattt 3120
gtaactgcct ctgttttagg agtataagta ttacttcctt gtgggtctatt gtgaagtaaa 3180
aagtagaccc ttgcatatac tattcttggt tgtgttcac ttaatgtttt tgtacagcta 3240
aatcaaatgt aatttataga gttagtttca tcaacctaat gaatgctagt taaatttgaa 3300
ttccttgga tttatcgtat attgtattca ctgagattat gaagggacaa atgttaatct 3360
tttgtttcca gaaaaagttg ggctttccca agcagttcta ttacccggtt cagaattgct 3420
tcatccaaaa atcatctgat ggtatagatg gacccctagc cttttcatta cctgatggta 3480
gaaataaaat aattgatttt a 3501

```

<210> 6

<211> 752

<212> PRT

<213> Homo sapiens

<400> 6

```

Met Arg Leu Leu Gly Ala Ala Ala Val Ala Ala Leu Gly Arg Gly Arg
  1              5              10              15

Ala Pro Ala Ser Leu Gly Trp Gln Arg Lys Gln Val Asn Trp Lys Ala
      20              25              30

Cys Arg Trp Ser Ser Ser Gly Val Ile Pro Asn Glu Lys Ile Arg Asn
      35              40              45

Ile Gly Ile Ser Ala His Ile Asp Ser Gly Lys Thr Thr Leu Thr Glu
      50              55              60

Arg Val Leu Tyr Tyr Thr Gly Arg Ile Ala Lys Met His Glu Val Lys
      65              70              75              80

Gly Lys Asp Gly Val Gly Ala Val Met Asp Ser Met Glu Leu Glu Arg
      85              90              95

Gln Arg Gly Ile Thr Ile Gln Ser Ala Ala Thr Phe Thr Met Trp Lys
      100             105             110

Asp Val Asn Ile Asn Ile Ile Asp Thr Pro Gly His Val Asp Phe Thr
      115             120             125

Ile Glu Val Glu Arg Ala Leu Arg Val Leu Asp Gly Ala Val Leu Val
      130             135             140

Leu Cys Ala Val Gly Gly Val Gln Cys Gln Thr Met Thr Val Asn Arg
      145             150             155             160

Gln Met Lys Arg Tyr Asn Val Pro Phe Leu Thr Phe Ile Asn Lys Leu
      165             170             175

Asp Arg Met Gly Ser Asn Pro Ala Arg Ala Leu Gln Gln Met Arg Ser
      180             185             190

Lys Leu Asn His Asn Thr Ala Phe Met Gln Ile Pro Met Gly Leu Glu
      195             200             205

Gly Asn Phe Lys Gly Ile Val Asp Leu Ile Glu Glu Arg Ala Ile Tyr
      210             215             220

```


Phe	Asp	Gly	Asp	Phe	Ser	Gln	Ile	Val	Arg	Tyr	Gly	Glu	Ile	Pro	Ala	
225					230					235					240	
Glu	Leu	Arg	Ala	Ala	Ala	Thr	Asp	His	Arg	Gln	Glu	Leu	Ile	Glu	Cys	
			245						250					255		
Val	Ala	Asn	Ser	Asp	Glu	Gln	Leu	Gly	Glu	Met	Phe	Leu	Glu	Glu	Lys	
			260					265					270			
Ile	Pro	Ser	Ile	Ser	Asp	Leu	Lys	Leu	Ala	Ile	Arg	Arg	Ala	Thr	Leu	
		275					280					285				
Lys	Arg	Ser	Phe	Thr	Pro	Val	Phe	Leu	Gly	Ser	Ala	Leu	Lys	Asn	Lys	
	290					295					300					
Gly	Val	Gln	Pro	Leu	Leu	Asp	Ala	Val	Leu	Glu	Tyr	Leu	Pro	Asn	Pro	
305					310					315					320	
Ser	Glu	Val	Gln	Asn	Tyr	Ala	Ile	Leu	Asn	Lys	Lys	Asp	Asp	Ser	Lys	
				325					330						335	
Glu	Lys	Thr	Lys	Ile	Leu	Met	Asn	Ser	Ser	Arg	His	Asn	Ser	His	Pro	
			340					345					350			
Phe	Val	Gly	Leu	Ala	Phe	Pro	Leu	Glu	Val	Gly	Arg	Phe	Gly	Gln	Leu	
		355					360					365				
Thr	Tyr	Val	Arg	Ser	Tyr	Gln	Gly	Glu	Leu	Lys	Lys	Gly	Asp	Thr	Ile	
		370				375						380				
Tyr	Asn	Thr	Arg	Thr	Arg	Lys	Lys	Val	Arg	Leu	Gln	Arg	Leu	Ala	Arg	
385					390					395					400	
Met	His	Ala	Asp	Met	Met	Glu	Ala	Ser	Thr	Glu	Glu	Val	Tyr	Ala	Gly	
				405					410					415		
Asp	Ile	Cys	Ala	Leu	Phe	Gly	Ile	Asp	Cys	Ala	Ser	Gly	Asp	Thr	Phe	
			420					425					430			
Thr	Asp	Lys	Ala	Asn	Ser	Gly	Leu	Ser	Met	Glu	Ser	Ile	His	Val	Pro	
		435					440					445				
Asp	Pro	Val	Ile	Ser	Ile	Ala	Met	Lys	Pro	Ser	Asn	Lys	Asn	Asp	Leu	
	450					455					460					
Glu	Lys	Phe	Ser	Lys	Gly	Ile	Gly	Arg	Phe	Thr	Arg	Glu	Asp	Pro	Thr	
465					470					475					480	
Phe	Lys	Val	Tyr	Phe	Asp	Thr	Glu	Asn	Lys	Glu	Thr	Val	Ile	Ser	Gly	
				485					490					495		
Met	Gly	Glu	Leu	His	Leu	Glu	Ile	Tyr	Ala	Gln	Arg	Leu	Glu	Arg	Glu	
			500					505					510			
Tyr	Gly	Cys	Pro	Cys	Ile	Thr	Gly	Lys	Pro	Lys	Val	Ala	Phe	Arg	Glu	
		515					520					525				

Thr Ile Thr Ala Pro Val Pro Phe Asp Phe Thr His Lys Lys Gln Ser
 530 535 540
 Gly Gly Ala Gly Gln Tyr Gly Lys Val Ile Gly Val Leu Glu Pro Leu
 545 550 555 560
 Asp Pro Glu Asp Tyr Thr Lys Leu Glu Phe Ser Asp Glu Thr Phe Gly
 565 570 575
 Ser Asn Ile Pro Lys Gln Phe Val Pro Ala Val Glu Lys Gly Phe Leu
 580 585 590
 Asp Ala Cys Glu Lys Gly Pro Leu Ser Gly His Lys Leu Ser Gly Leu
 595 600 605
 Arg Phe Val Leu Gln Asp Gly Ala His His Met Val Asp Ser Asn Glu
 610 615 620
 Ile Ser Phe Ile Arg Ala Gly Glu Gly Ala Leu Lys Gln Ala Leu Ala
 625 630 635 640
 Asn Ala Thr Leu Cys Ile Leu Glu Pro Ile Met Ala Val Glu Val Val
 645 650 655
 Ala Pro Asn Glu Phe Gln Gly Gln Val Ile Ala Gly Ile Asn Arg Arg
 660 665 670
 His Gly Val Ile Thr Gly Gln Asp Gly Val Glu Asp Tyr Phe Thr Leu
 675 680 685
 Tyr Ala Asp Val Pro Leu Asn Asp Met Phe Gly Tyr Ser Thr Glu Leu
 690 695 700
 Arg Ser Cys Thr Glu Gly Lys Gly Glu Tyr Thr Met Glu Tyr Ser Arg
 705 710 715 720
 Tyr Gln Pro Cys Leu Pro Ser Thr Gln Glu Asp Val Ile Asn Lys Tyr
 725 730 735
 Leu Glu Ala Thr Gly Gln Leu Pro Val Lys Lys Gly Lys Ala Lys Asn
 740 745 750

<210> 7

<211> 7506

<212> DNA

<213> Homo sapiens

<400> 7

```

gccgcgggag caggcggagg cggaggcggc gggggcagga ggatgtcgca gccgccgtg 60
ctccccgcct cggcggagac tcggaagttc acccgggcgc tgagtaagcc gggcacggcg 120
gccgagctgc ggcagagcgt gtctgaggtg gtgcgcggct ccgtgctcct ggcaaagcca 180
aagctaattg agccactcga ctatgaaaat gtcacgtcc agaagaagac tcagatcctg 240
aacgactgtt tacgggagat gctgctcttc ccttacgatg actttcagac ggccatcctg 300
agacgacagg gtcgatacat atgctcaaca gtgcctgcga aggcggaaga ggaagcacag 360
agcttgtttg ttacagagtg catcaaaacc tataactctg actggcatct tgtgaactat 420
aaatatgaag attactcagg agagtttcga cagcttccga acaaagtggg caagttggat 480
aaacttccag ttcatgtcta tgaagttgac gaggagggtcg acaaagatga ggatgctgcc 540

```

tcccttggtt	cccagaaagg	tgggatcacc	aagcatggct	ggctgtacaa	aggcaacatg	600
aacagtgcc	tcagcgtgac	catgaggtca	tttaagagac	gatttttcca	cctgattcaa	660
cttggcgatg	gacctataa	atttgaattt	ttaaaagatc	tccaaaagga	accaaagga	720
tcaatatttc	tgggattcct	gtatgggggtg	tcgttcagga	acaacaaagt	caggcgtttt	780
gcttttgagc	tcaagatgca	ggacaaaagt	agttatctct	tggcagcaga	cagtgaagtg	840
gaaatggaag	aatggatcac	aattctaaat	aagatcctcc	agctcaactt	tgaagctgca	900
atgcaagaaa	agcgaaatgg	cgactctcac	gaagatgatg	aacaaagcaa	attggaaggt	960
tctgggttccg	gtttagatag	ctacctgccg	gaacttgcca	agagtgcaag	agaagcagaa	1020
atcaaaactga	aaagtgaaaag	cagagtcaaaa	cttttttatt	tggaccacaga	tgcccagaag	1080
cttgacttct	catcagctga	gccagaagtg	aagtcatttg	aagagaagtt	tggaaaaagg	1140
atccttgatc	agtgcaatga	tttatctttt	aatttgcaat	gctgtgttgc	cgaaaatgaa	1200
gaaggaccca	ctacaaatgt	tgaacctttc	tttgttactc	tatccctggt	tgacataaaa	1260
tacaaccgga	agatttctgc	cgatttccac	gtagacctga	accatttctc	agtgaggcaa	1320
atgatcgcca	ccacgtcccc	ggcgctgatg	aatggcagtg	ggccggaac	ccaatctgcc	1380
ctcaggggca	tcttcatga	agccgccatg	cagtatccga	agcagggaa	attttcagtc	1440
acttgctctc	atccagatat	atctcttggt	gccagaattg	aaaaagtcct	tcaggggagc	1500
atcacacatt	gcgctgagcc	atatatgaaa	agttcagact	cttctaaggt	ggcccagaag	1560
gtgctgaaga	atgccaaagca	ggcatgcca	agactaggac	agtatagaat	gccatttgct	1620
tgggcagcaa	ggacattggt	taaggatgca	tctggaaatc	ttgacaaaa	tgccagattt	1680
tctgccatct	acaggcaaga	cagcaataag	ctatccaatg	atgacatgct	caagttactt	1740
gcaggtttc	ggaacactga	gaagatggct	aagctcccag	tgattttagg	caactagac	1800
attacaattg	ataatgtttc	ctcagacttc	cctaattatg	ttaattcatc	atacattccc	1860
acaaaacaat	ttgaaacctg	cagtaaaaact	cccatcacgt	ttgaagtgga	ggaatttggt	1920
ccctgcatac	caaaacacac	tcagccttac	accatctaca	ccaatcacct	ttacgtttat	1980
cctaagtact	tgaatacga	cagtcagaag	tcttttgcca	aggctagaaa	tattgcgatt	2040
tgcattgaat	tcaaagattc	agatgaggaa	gactctcagc	cccttaagtg	catttatggc	2100
agacctgggtg	ggccagtttt	cacaagaagc	gcctttgctg	cagttttaca	ccatcaccaa	2160
aacccagaat	tttatgatga	gattaaaata	gagttgccca	ctcagctgca	tgaaaagcac	2220
cacctgttgc	tcacattctt	ccatgtcagc	tgtagaact	caagtaaagg	aagcacgaag	2280
aagagggatg	tcgttgaaac	ccaagtggc	tactcctggc	ttccctcct	gaaagacgga	2340
aggggtgggtg	caagcgagca	gcacatccc	gtctcggcga	accttccttc	gggtatctt	2400
ggctaccagg	agcttgggat	gggcaggcat	tatggtccgg	aaattaaatg	ggtagatgga	2460
ggcaagccac	tgttgaaaat	ttccactcat	ctggtttcta	cagtgtatac	tcaggatcag	2520
catttacata	attttttcca	gtactgtcag	aaaaccgaat	ctggagccca	agccttagga	2580
aacgaacttg	taaagtacct	taagagtctg	catgcgatgg	aaggccacgt	gatgatcgcc	2640
ttcttgccca	ctatcctaaa	ccagctgttc	cgagtcctca	ccagagccac	acaggaagaa	2700
gtcgcggtta	acgtgactcg	ggtcattatt	catgtgggtg	cccagtgcca	tgaggaagga	2760
ttggagagcc	acttgagggtc	atatgttaag	tacgcgtata	aggctgagcc	atatgttgcc	2820
tctgaatata	agacagtga	tgaagaactg	accaaattcca	tgaccacgat	tctcaagcct	2880
tctgccgatt	tcctcaccag	caacaaacta	ctgaagtact	catgggtttt	ctttgatgta	2940
ctgatcaaat	ctatggctca	gcatttgata	gagaactcca	aagttaagtt	gctgcgaaac	3000
cagagatttc	ctgcaccta	tcacatgca	gtggaaaccg	ttgtaaatat	gctgatgcca	3060
cacatcactc	agaagtttcg	agataatcca	gaggcatcta	agaacgcgaa	tcatagcctt	3120
gctgtcttca	tcaagagatg	tttcaccttc	atggacaggg	gctttgtctt	caagcagatc	3180
aacaactaca	ttagctgttt	tgctcctgga	gacccaaaga	ccctctttga	atacaagttt	3240
gaattttctcc	gtgtagtgtg	caaccatgaa	cattatattc	cgttgaactt	accaatgcca	3300
tttgaaaaag	gcaggattca	aagataccaa	gacctccagc	ttgactactc	attaacagat	3360
gagttctgca	gaaaccactt	cttgggtggg	ctgttactga	gggaggtggg	gacagccctc	3420
caggagttcc	gggaggtccg	tctgatcgcc	atcagtgtgc	tcaagaacct	gctgataaag	3480
cattcttttg	atgacagata	tgcttcaagg	agccatcagg	caaggatagc	cacctctac	3540
ctgcctctgt	ttggtctgct	gattgaaaac	gtccagcgga	tcaatgtgag	ggatgtgtca	3600
cccttccctg	tgaacgcggg	catgactgtg	aaggatgaat	ccctggctct	accagctgtg	3660
aatccgctgg	tgacgccgca	gaagggaagc	accctggaca	acagcctgca	caaggacctg	3720
ctgggcgcca	tctccggcat	tgcttctcca	tatacaacct	caactccaaa	catcaacagt	3780
gtgagaaatg	ctgattcgag	aggatctctc	ataagcacag	attcgggtaa	cagccttcca	3840
gaaaggaata	gtgagaagag	caattccctg	gataagcacc	aacaaagtag	cacattggga	3900
aattccgtgg	ttcgctgtga	ttaacttgac	cagcttgaga	ttaagagcct	actgatgtgt	3960
ttcctctaca	tcttaagag	catgtctgat	gatgctttgt	ttacatattg	gaacaaggct	4020

tcaacatctg	aacttatgga	tttttttaca	atatctgaag	tctgcctgca	ccagttccag	4080
tacatgggga	agcgatacat	agccagaaca	ggaatgatgc	atgccagatt	gcagcagctg	4140
ggcagcctgg	ataactctct	cactttttaac	cacagctatg	gccactcgga	cgcatagtgt	4200
ctgcaccagt	cattacttga	agccaacatt	gctactgagg	tttgcctgac	agctctggac	4260
acgcttttct	tatttacatt	ggcgtttaag	aaccagctcc	tggccgacca	tggacataat	4320
cctctcatga	aaaaagtttt	tgatgtctac	ctgtgttttc	ttcaaaaaca	tcagtctgaa	4380
acggctttta	aaaatgtctt	cactgcctta	aggtccttaa	tttataagtt	tccctcaaca	4440
ttctatgaag	ggagagcgga	catgtgtgcg	gctctgtgtt	acgagattct	caagtgtctg	4500
aactccaagc	tgagctccat	caggacggag	gcctcccagc	tgctctactt	cctgatgagg	4560
aacaactttg	attacactgg	aaagaagtcc	tttgtccgga	cacatttgca	agtcatacata	4620
tctgtcagcc	agctgatagc	agacgttggt	ggcattgggg	gaaccagatt	ccagcagctc	4680
ctgtccatga	tcaacaactg	tgccaacagt	gaccggctta	ttaagcacac	cagcttctct	4740
tctgatgtga	aggacttaac	caaaaggata	cgcacggtgc	taatggccac	cgcccagatg	4800
aaggagcatg	agaacgaccc	agagatgctg	gtggacctcc	agtacagcct	ggccaaatcc	4860
tatgccagca	cgcccagagc	caggaagacg	tggctcgaca	gcatggccag	gatccatgtc	4920
aaaaatggcg	atctctcaga	ggcagcaatg	tgctatgtcc	acgtaacagc	cctagtggca	4980
gaatatctca	cacggaaaga	agcagtccag	tgggagccgc	cccttctccc	ccacagccat	5040
agcgcctgcc	tgaggaggag	ccggggaggc	gtgttttagac	aaggatgcac	cgccttcagg	5100
gtcattaccc	caaacatcga	cgaggaggcc	tccatgatgg	aagacgtggg	gatgcaggat	5160
gtccatttca	acgaggatgt	gctgatggag	ctccttgagc	agtgcgcaga	tggactctgg	5220
aaagccgagc	gctacgagct	cattgcccag	atctacaaac	ttatcatccc	cattttatgag	5280
aagcggaggg	attttgagag	gctggcccat	ctgtatgaca	cgctgcaccg	ggcctacagc	5340
aaagtgaccg	aggtcatgca	ctcggggccg	aggcttctgg	ggacctactt	ccgggtagcc	5400
ttcttcgggc	aggcagcgca	ataccagttt	acagacagtg	aaacagatgt	ggagggattc	5460
tttgaagatg	aagatggaaa	ggagtataat	tacaaggaac	ccaaactcac	accgctgtcg	5520
gaaatttctc	agagactcct	taaactgtac	tcggataaat	ttggttctga	aaatgtcaaa	5580
atgatacagg	attctggcaa	ggtcaaccct	aaggatctgg	attctaagta	tgcctacatc	5640
caggtgactc	acgtcatccc	cttctttgac	gaaaaagagt	tgcaagaaag	gaaaacagag	5700
tttgagagat	cccacaacat	ccgccgcttc	atgtttgaga	tgccatttac	gcagaccggg	5760
aagaggcagg	gcgggggtga	agagcagtg	aaacggcgca	ccatcctgac	agccatacac	5820
tgcttccctt	atgtgaagaa	gcgcattccc	gtcatgtacc	agcaccacac	tgacctgaac	5880
cccatcgagg	tggccattga	cgagatgagt	aagaagggtg	cggagctccg	gcagctgtgc	5940
tcctcggccg	aggtggacat	gatcaaaact	cagctcaaac	tccagggcag	cgtgagtgtt	6000
caggtcaatg	ctggcccact	agcatatg	cgagctttct	tagatgatac	aaacacaaag	6060
cgatatcctg	acaataaagt	gaagctgctt	aaggaaagtt	tcaggcaatt	tgtggaagct	6120
tgcggccaag	ccttagcggt	aaacgaacgt	ctgattaaag	aagaccagct	cgagtatcag	6180
gaagaaatga	aagccaacta	cagggaaatg	gcgaaggagc	tttctgaaat	catgcatgag	6240
cagatctgcc	ccctggagga	gaagacgagc	gtcttaccga	attcccttca	catcttcaac	6300
gccatcagtg	ggactccaac	agcacaaatg	gttcacggga	tgaccagctc	gtcttcggct	6360
gtgtgattac	atctcatggc	ccgtgtgtgg	ggactgtgct	tgtcatttgc	aaactcagga	6420
tgctttccaa	agccaatcac	tggggagacc	gagcacaggg	aggaccaagg	ggaaggggag	6480
agaaaggaaa	taaagaacaa	cgttatttct	taacagactt	tctataggag	ttgtaagaag	6540
gtgcacatat	tttttttaaat	ctcactggca	atattcaaa	ttttcattgt	gtcttaacaa	6600
aggtgtggta	gacactcttg	agctggactt	agattttatt	cttccttgca	gagtgtgtt	6660
agaatagatg	gcctacagaa	aaaaaagggt	ctgggatcta	catggcaggg	agggctgcac	6720
tgacattgat	gcctggggga	ccttttgcc	cgaggctgag	ctggaaaatc	ttgaaaatat	6780
tttttttttc	ctgtggcaca	ttcagggtga	atacaagaac	tatttttgtg	actagttttt	6840
gatgacctaa	gggaactgac	cattgtaatt	tttgtaccag	tgaaccagga	gatttagtgc	6900
ttttatatct	atttccttgc	atttaagaaa	atatgaaagc	ttaagggaatt	atgtgagctt	6960
aaaactagtc	aagcagttta	gaaccaaagg	cctatatata	taaccgcaac	tatgctgaaa	7020
agtacaaagt	agtacagtat	attgttatgt	acatatcatt	gttaatacac	tcctggcatt	7080
ctgtacatat	atgtattaca	tttctacatt	tttaatactc	acatgggctt	atgcattaag	7140
tttaattgtg	ataaatttgt	gctgttccag	tatatgcaat	acactttaat	gttttattct	7200
tgtacataaa	aatgtgcaat	atggagatgt	atacagctct	tactatatta	ggtttataaa	7260
cagttttaag	aatttcatcc	ttttgccaaa	atgggtggag	atgtaattgg	ttaatcataa	7320
atcctgtggg	gaatgggtgg	gtactttaaa	gctgtcacca	tggttatatt	tcttttaaga	7380
ctttaattta	gtaattttat	atttgggaaa	ataaagggtt	ttaattttat	ttaactggaa	7440
tcactgcctc	gctgtaatta	aacattctgt	accacatctg	tattaaaaag	acattgctga	7500

ccatta

<210> 8
 <211> 2107
 <212> PRT
 <213> Homo sapiens

<400> 8
 Met Ser Gln Pro Pro Leu Leu Pro Ala Ser Ala Glu Thr Arg Lys Phe
 1 5 10 15
 Thr Arg Ala Leu Ser Lys Pro Gly Thr Ala Ala Glu Leu Arg Gln Ser
 20 25 30
 Val Ser Glu Val Val Arg Gly Ser Val Leu Leu Ala Lys Pro Lys Leu
 35 40 45
 Ile Glu Pro Leu Asp Tyr Glu Asn Val Ile Val Gln Lys Lys Thr Gln
 50 55 60
 Ile Leu Asn Asp Cys Leu Arg Glu Met Leu Leu Phe Pro Tyr Asp Asp
 65 70 75 80
 Phe Gln Thr Ala Ile Leu Arg Arg Gln Gly Arg Tyr Ile Cys Ser Thr
 85 90 95
 Val Pro Ala Lys Ala Glu Glu Glu Ala Gln Ser Leu Phe Val Thr Glu
 100 105 110
 Cys Ile Lys Thr Tyr Asn Ser Asp Trp His Leu Val Asn Tyr Lys Tyr
 115 120 125
 Glu Asp Tyr Ser Gly Glu Phe Arg Gln Leu Pro Asn Lys Val Val Lys
 130 135 140
 Leu Asp Lys Leu Pro Val His Val Tyr Glu Val Asp Glu Glu Val Asp
 145 150 155 160
 Lys Asp Glu Asp Ala Ala Ser Leu Gly Ser Gln Lys Gly Gly Ile Thr
 165 170 175
 Lys His Gly Trp Leu Tyr Lys Gly Asn Met Asn Ser Ala Ile Ser Val
 180 185 190
 Thr Met Arg Ser Phe Lys Arg Arg Phe Phe His Leu Ile Gln Leu Gly
 195 200 205
 Asp Gly Ser Tyr Lys Phe Glu Phe Leu Lys Asp Leu Gln Lys Glu Pro
 210 215 220
 Lys Gly Ser Ile Phe Leu Gly Phe Leu Tyr Gly Val Ser Phe Arg Asn
 225 230 235 240
 Asn Lys Val Arg Arg Phe Ala Phe Glu Leu Lys Met Gln Asp Lys Ser
 245 250 255

Ser Tyr Leu Leu Ala Ala Asp Ser Glu Val Glu Met Glu Glu Trp Ile
 260 265 270
 Thr Ile Leu Asn Lys Ile Leu Gln Leu Asn Phe Glu Ala Ala Met Gln
 275 280 285
 Glu Lys Arg Asn Gly Asp Ser His Glu Asp Asp Glu Gln Ser Lys Leu
 290 295 300
 Glu Gly Ser Gly Ser Gly Leu Asp Ser Tyr Leu Pro Glu Leu Ala Lys
 305 310 315 320
 Ser Ala Arg Glu Ala Glu Ile Lys Leu Lys Ser Glu Ser Arg Val Lys
 325 330 335
 Leu Phe Tyr Leu Asp Pro Asp Ala Gln Lys Leu Asp Phe Ser Ser Ala
 340 345 350
 Glu Pro Glu Val Lys Ser Phe Glu Glu Lys Phe Gly Lys Arg Ile Leu
 355 360 365
 Val Lys Cys Asn Asp Leu Ser Phe Asn Leu Gln Cys Cys Val Ala Glu
 370 375 380
 Asn Glu Glu Gly Pro Thr Thr Asn Val Glu Pro Phe Phe Val Thr Leu
 385 390 395 400
 Ser Leu Phe Asp Ile Lys Tyr Asn Arg Lys Ile Ser Ala Asp Phe His
 405 410 415
 Val Asp Leu Asn His Phe Ser Val Arg Gln Met Ile Ala Thr Thr Ser
 420 425 430
 Pro Ala Leu Met Asn Gly Ser Gly Pro Glu Thr Gln Ser Ala Leu Arg
 435 440 445
 Gly Ile Leu His Glu Ala Ala Met Gln Tyr Pro Lys Gln Gly Ile Phe
 450 455 460
 Ser Val Thr Cys Pro His Pro Asp Ile Phe Leu Val Ala Arg Ile Glu
 465 470 475 480
 Lys Val Leu Gln Gly Ser Ile Thr His Cys Ala Glu Pro Tyr Met Lys
 485 490 495
 Ser Ser Asp Ser Ser Lys Val Ala Gln Lys Val Leu Lys Asn Ala Lys
 500 505 510
 Gln Ala Cys Gln Arg Leu Gly Gln Tyr Arg Met Pro Phe Ala Trp Ala
 515 520 525
 Ala Arg Thr Leu Phe Lys Asp Ala Ser Gly Asn Leu Asp Lys Asn Ala
 530 535 540
 Arg Phe Ser Ala Ile Tyr Arg Gln Asp Ser Asn Lys Leu Ser Asn Asp
 545 550 555 560

Asp Met Leu Lys Leu Leu Ala Asp Phe Arg Lys Pro Glu Lys Met Ala
 565 570 575
 Lys Leu Pro Val Ile Leu Gly Asn Leu Asp Ile Thr Ile Asp Asn Val
 580 585 590
 Ser Ser Asp Phe Pro Asn Tyr Val Asn Ser Ser Tyr Ile Pro Thr Lys
 595 600 605
 Gln Phe Glu Thr Cys Ser Lys Thr Pro Ile Thr Phe Glu Val Glu Glu
 610 615 620
 Phe Val Pro Cys Ile Pro Lys His Thr Gln Pro Tyr Thr Ile Tyr Thr
 625 630 635 640
 Asn His Leu Tyr Val Tyr Pro Lys Tyr Leu Lys Tyr Asp Ser Gln Lys
 645 650 655
 Ser Phe Ala Lys Ala Arg Asn Ile Ala Ile Cys Ile Glu Phe Lys Asp
 660 665 670
 Ser Asp Glu Glu Asp Ser Gln Pro Leu Lys Cys Ile Tyr Gly Arg Pro
 675 680 685
 Gly Gly Pro Val Phe Thr Arg Ser Ala Phe Ala Ala Val Leu His His
 690 695 700
 His Gln Asn Pro Glu Phe Tyr Asp Glu Ile Lys Ile Glu Leu Pro Thr
 705 710 715 720
 Gln Leu His Glu Lys His His Leu Leu Leu Thr Phe Phe His Val Ser
 725 730 735
 Cys Asp Asn Ser Ser Lys Gly Ser Thr Lys Lys Arg Asp Val Val Glu
 740 745 750
 Thr Gln Val Gly Tyr Ser Trp Leu Pro Leu Leu Lys Asp Gly Arg Val
 755 760 765
 Val Thr Ser Glu Gln His Ile Pro Val Ser Ala Asn Leu Pro Ser Gly
 770 775 780
 Tyr Leu Gly Tyr Gln Glu Leu Gly Met Gly Arg His Tyr Gly Pro Glu
 785 790 795 800
 Ile Lys Trp Val Asp Gly Gly Lys Pro Leu Leu Lys Ile Ser Thr His
 805 810 815
 Leu Val Ser Thr Val Tyr Thr Gln Asp Gln His Leu His Asn Phe Phe
 820 825 830
 Gln Tyr Cys Gln Lys Thr Glu Ser Gly Ala Gln Ala Leu Gly Asn Glu
 835 840 845
 Leu Val Lys Tyr Leu Lys Ser Leu His Ala Met Glu Gly His Val Met
 850 855 860

Ile Ala Phe Leu Pro Thr Ile Leu Asn Gln Leu Phe Arg Val Leu Thr
 865 870 875 880
 Arg Ala Thr Gln Glu Glu Val Ala Val Asn Val Thr Arg Val Ile Ile
 885 890 895
 His Val Val Ala Gln Cys His Glu Glu Gly Leu Glu Ser His Leu Arg
 900 905 910
 Ser Tyr Val Lys Tyr Ala Tyr Lys Ala Glu Pro Tyr Val Ala Ser Glu
 915 920 925
 Tyr Lys Thr Val His Glu Glu Leu Thr Lys Ser Met Thr Thr Ile Leu
 930 935 940
 Lys Pro Ser Ala Asp Phe Leu Thr Ser Asn Lys Leu Leu Lys Tyr Ser
 945 950 955 960
 Trp Phe Phe Phe Asp Val Leu Ile Lys Ser Met Ala Gln His Leu Ile
 965 970 975
 Glu Asn Ser Lys Val Lys Leu Leu Arg Asn Gln Arg Phe Pro Ala Ser
 980 985 990
 Tyr His His Ala Val Glu Thr Val Val Asn Met Leu Met Pro His Ile
 995 1000 1005
 Thr Gln Lys Phe Arg Asp Asn Pro Glu Ala Ser Lys Asn Ala Asn His
 1010 1015 1020
 Ser Leu Ala Val Phe Ile Lys Arg Cys Phe Thr Phe Met Asp Arg Gly
 1025 1030 1035 1040
 Phe Val Phe Lys Gln Ile Asn Asn Tyr Ile Ser Cys Phe Ala Pro Gly
 1045 1050 1055
 Asp Pro Lys Thr Leu Phe Glu Tyr Lys Phe Glu Phe Leu Arg Val Val
 1060 1065 1070
 Cys Asn His Glu His Tyr Ile Pro Leu Asn Leu Pro Met Pro Phe Gly
 1075 1080 1085
 Lys Gly Arg Ile Gln Arg Tyr Gln Asp Leu Gln Leu Asp Tyr Ser Leu
 1090 1095 1100
 Thr Asp Glu Phe Cys Arg Asn His Phe Leu Val Gly Leu Leu Leu Arg
 1105 1110 1115 1120
 Glu Val Gly Thr Ala Leu Gln Glu Phe Arg Glu Val Arg Leu Ile Ala
 1125 1130 1135
 Ile Ser Val Leu Lys Asn Leu Leu Ile Lys His Ser Phe Asp Asp Arg
 1140 1145 1150
 Tyr Ala Ser Arg Ser His Gln Ala Arg Ile Ala Thr Leu Tyr Leu Pro
 1155 1160 1165

Leu Phe Gly Leu Leu Ile Glu Asn Val Gln Arg Ile Asn Val Arg Asp
 1170 1175 1180
 Val Ser Pro Phe Pro Val Asn Ala Gly Met Thr Val Lys Asp Glu Ser
 1185 1190 1195 1200
 Leu Ala Leu Pro Ala Val Asn Pro Leu Val Thr Pro Gln Lys Gly Ser
 1205 1210 1215
 Thr Leu Asp Asn Ser Leu His Lys Asp Leu Leu Gly Ala Ile Ser Gly
 1220 1225 1230
 Ile Ala Ser Pro Tyr Thr Thr Ser Thr Pro Asn Ile Asn Ser Val Arg
 1235 1240 1245
 Asn Ala Asp Ser Arg Gly Ser Leu Ile Ser Thr Asp Ser Gly Asn Ser
 1250 1255 1260
 Leu Pro Glu Arg Asn Ser Glu Lys Ser Asn Ser Leu Asp Lys His Gln
 1265 1270 1275 1280
 Gln Ser Ser Thr Leu Gly Asn Ser Val Val Arg Cys Asp Lys Leu Asp
 1285 1290 1295
 Gln Ser Glu Ile Lys Ser Leu Leu Met Cys Phe Leu Tyr Ile Leu Lys
 1300 1305 1310
 Ser Met Ser Asp Asp Ala Leu Phe Thr Tyr Trp Asn Lys Ala Ser Thr
 1315 1320 1325
 Ser Glu Leu Met Asp Phe Phe Thr Ile Ser Glu Val Cys Leu His Gln
 1330 1335 1340
 Phe Gln Tyr Met Gly Lys Arg Tyr Ile Ala Arg Thr Gly Met Met His
 1345 1350 1355 1360
 Ala Arg Leu Gln Gln Leu Gly Ser Leu Asp Asn Ser Leu Thr Phe Asn
 1365 1370 1375
 His Ser Tyr Gly His Ser Asp Ala Asp Val Leu His Gln Ser Leu Leu
 1380 1385 1390
 Glu Ala Asn Ile Ala Thr Glu Val Cys Leu Thr Ala Leu Asp Thr Leu
 1395 1400 1405
 Ser Leu Phe Thr Leu Ala Phe Lys Asn Gln Leu Leu Ala Asp His Gly
 1410 1415 1420
 His Asn Pro Leu Met Lys Lys Val Phe Asp Val Tyr Leu Cys Phe Leu
 1425 1430 1435 1440
 Gln Lys His Gln Ser Glu Thr Ala Leu Lys Asn Val Phe Thr Ala Leu
 1445 1450 1455
 Arg Ser Leu Ile Tyr Lys Phe Pro Ser Thr Phe Tyr Glu Gly Arg Ala
 1460 1465 1470

Asp Met Cys Ala Ala Leu Cys Tyr Glu Ile Leu Lys Cys Cys Asn Ser
 1475 1480 1485
 Lys Leu Ser Ser Ile Arg Thr Glu Ala Ser Gln Leu Leu Tyr Phe Leu
 1490 1495 1500
 Met Arg Asn Asn Phe Asp Tyr Thr Gly Lys Lys Ser Phe Val Arg Thr
 1505 1510 1515 1520
 His Leu Gln Val Ile Ile Ser Val Ser Gln Leu Ile Ala Asp Val Val
 1525 1530 1535
 Gly Ile Gly Gly Thr Arg Phe Gln Gln Ser Leu Ser Ile Ile Asn Asn
 1540 1545 1550
 Cys Ala Asn Ser Asp Arg Leu Ile Lys His Thr Ser Phe Ser Ser Asp
 1555 1560 1565
 Val Lys Asp Leu Thr Lys Arg Ile Arg Thr Val Leu Met Ala Thr Ala
 1570 1575 1580
 Gln Met Lys Glu His Glu Asn Asp Pro Glu Met Leu Val Asp Leu Gln
 1585 1590 1595 1600
 Tyr Ser Leu Ala Lys Ser Tyr Ala Ser Thr Pro Glu Leu Arg Lys Thr
 1605 1610 1615
 Trp Leu Asp Ser Met Ala Arg Ile His Val Lys Asn Gly Asp Leu Ser
 1620 1625 1630
 Glu Ala Ala Met Cys Tyr Val His Val Thr Ala Leu Val Ala Glu Tyr
 1635 1640 1645
 Leu Thr Arg Lys Glu Ala Val Gln Trp Glu Pro Pro Leu Leu Pro His
 1650 1655 1660
 Ser His Ser Ala Cys Leu Arg Arg Ser Arg Gly Gly Val Phe Arg Gln
 1665 1670 1675 1680
 Gly Cys Thr Ala Phe Arg Val Ile Thr Pro Asn Ile Asp Glu Glu Ala
 1685 1690 1695
 Ser Met Met Glu Asp Val Gly Met Gln Asp Val His Phe Asn Glu Asp
 1700 1705 1710
 Val Leu Met Glu Leu Leu Glu Gln Cys Ala Asp Gly Leu Trp Lys Ala
 1715 1720 1725
 Glu Arg Tyr Glu Leu Ile Ala Asp Ile Tyr Lys Leu Ile Ile Pro Ile
 1730 1735 1740
 Tyr Glu Lys Arg Arg Asp Phe Glu Arg Leu Ala His Leu Tyr Asp Thr
 1745 1750 1755 1760
 Leu His Arg Ala Tyr Ser Lys Val Thr Glu Val Met His Ser Gly Arg
 1765 1770 1775

Arg Leu Leu Gly Thr Tyr Phe Arg Val Ala Phe Phe Gly Gln Ala Ala
 1780 1785 1790
 Gln Tyr Gln Phe Thr Asp Ser Glu Thr Asp Val Glu Gly Phe Phe Glu
 1795 1800 1805
 Asp Glu Asp Gly Lys Glu Tyr Ile Tyr Lys Glu Pro Lys Leu Thr Pro
 1810 1815 1820
 Leu Ser Glu Ile Ser Gln Arg Leu Leu Lys Leu Tyr Ser Asp Lys Phe
 1825 1830 1835 1840
 Gly Ser Glu Asn Val Lys Met Ile Gln Asp Ser Gly Lys Val Asn Pro
 1845 1850 1855
 Lys Asp Leu Asp Ser Lys Tyr Ala Tyr Ile Gln Val Thr His Val Ile
 1860 1865 1870
 Pro Phe Phe Asp Glu Lys Glu Leu Gln Glu Arg Lys Thr Glu Phe Glu
 1875 1880 1885
 Arg Ser His Asn Ile Arg Arg Phe Met Phe Glu Met Pro Phe Thr Gln
 1890 1895 1900
 Thr Gly Lys Arg Gln Gly Gly Val Glu Glu Gln Cys Lys Arg Arg Thr
 1905 1910 1915 1920
 Ile Leu Thr Ala Ile His Cys Phe Pro Tyr Val Lys Lys Arg Ile Pro
 1925 1930 1935
 Val Met Tyr Gln His His Thr Asp Leu Asn Pro Ile Glu Val Ala Ile
 1940 1945 1950
 Asp Glu Met Ser Lys Lys Val Ala Glu Leu Arg Gln Leu Cys Ser Ser
 1955 1960 1965
 Ala Glu Val Asp Met Ile Lys Leu Gln Leu Lys Leu Gln Gly Ser Val
 1970 1975 1980
 Ser Val Gln Val Asn Ala Gly Pro Leu Ala Tyr Ala Arg Ala Phe Leu
 1985 1990 1995 2000
 Asp Asp Thr Asn Thr Lys Arg Tyr Pro Asp Asn Lys Val Lys Leu Leu
 2005 2010 2015
 Lys Glu Val Phe Arg Gln Phe Val Glu Ala Cys Gly Gln Ala Leu Ala
 2020 2025 2030
 Val Asn Glu Arg Leu Ile Lys Glu Asp Gln Leu Glu Tyr Gln Glu Glu
 2035 2040 2045
 Met Lys Ala Asn Tyr Arg Glu Met Ala Lys Glu Leu Ser Glu Ile Met
 2050 2055 2060
 His Glu Gln Ile Cys Pro Leu Glu Glu Lys Thr Ser Val Leu Pro Asn
 2065 2070 2075 2080

Ser Leu His Ile Phe Asn Ala Ile Ser Gly Thr Pro Thr Ser Thr Met
 2085 2090 2095

Val His Gly Met Thr Ser Ser Ser Ser Val Val
 2100 2105

<210> 9

<211> 6171

<212> DNA

<213> Homo sapiens

<400> 9

ttttgtttac	agggaaacacg	ggtctggctg	agagaaaatg	gccagcattt	tccaagtact	60
gtaaattcct	gtgcagaagg	catcgctgct	ttccggacag	actatggtca	ggtattcact	120
tacaagcaga	gcacaattac	ccaccagaag	gtgactgcta	tgcacccac	gaacgaggag	180
ggcgtggatg	acatggcgct	cttgacagag	ctccatggcg	gctccatcat	gtataactta	240
ttccagcggg	ataagagaaa	tcaaatatgg	acctacatcg	gctccatcct	ggcctctgtg	300
aaccctacc	agcccatcgc	cgggctgtac	gagcctgcc	ccatggagca	gtacagccgg	360
cgccacctgg	gcgagctgcc	cccgcacatc	ttcgccatcg	ccaacgagtg	ctaccgctgc	420
ctgtggaagc	gccacgacaa	ccagtgcac	ctcatcaagg	gtgaaagtgg	ggcaggtaaa	480
accgaaagca	ctaaattgat	cctcaagtgt	ctgtcagtc	tcagtcaaca	gtctttgga	540
ttgtccttaa	aggagaagac	atcctgtgtt	gaacgagcta	ttcttgaaag	cagcccatc	600
atggaagctt	tcggcaatgc	gaagaccgtg	tacaacaaca	actctagtcg	ctttggaag	660
ttgttcagc	tgaacatctg	tcagaaagga	aatattcagg	gcgggagaat	tgtagattgt	720
atcctctctt	cccagaaccg	agtagtaagg	caaaatccc	gggaaaggaa	ttatcacata	780
ttttatgcac	tgctggcagg	gctggaacat	gaagaaagag	aagaatttta	tttatctacg	840
ccagaaaact	accactactt	gaatcagctt	ggatgtgtag	aagacaagac	aatcagtgac	900
caggaattcct	ttaggggaagt	tattacggca	atggacgtga	tgcagttcag	caaggaggaa	960
gttcgggaag	tgctgaggct	gcttgctggt	atactgcatc	ttgggaacat	agaattttatc	1020
actgctggtg	gggcacagggt	ttccttcaaa	acagctttgg	gcagatctgc	ggagttactt	1080
gggctggacc	caacacagct	cacagatgct	ttgaccaga	gatcaatgtt	cctcagggga	1140
gaagagatcc	tcacgcctct	caatgttcaa	caggcagtag	acagcaggga	ctccctggcc	1200
atggctctgt	atgcgtgctg	ctttgagtgg	gtaatcaaga	agatcaacag	caggatcaaa	1260
ggcaatgagg	acttcaagtc	tattggcatc	ctcgacatct	ttggatttga	aaactttgag	1320
gttaatcact	ttgaacagtt	caatataaac	tatgcaaacy	agaaacttca	ggagtacttc	1380
aacaagcata	ttttttcttt	agaacaacta	gaatatagca	gggaaggatt	agtgtgggaa	1440
gatattgact	ggatagacaa	tggagaatgc	ctggacttga	ttgagaagaa	acttggcctc	1500
ctagccctta	tcaatgaaga	aagccatttt	cctcaagcca	cagacagcac	cttattggag	1560
aagctacaca	gtcagcatgc	gaataaccac	tttttgtga	agccagaggt	tgcagttaac	1620
aattttggag	tgaagcacta	tgctggagag	gtgcaatatg	atgtccgagg	tatcttggag	1680
aagaacagag	atacatttcg	agatgacctt	ctcaatttgc	taagagaaag	ccggtttgac	1740
tttatctacg	atctttttga	acatgtttca	agccgcaaca	accaggatac	cttgaaatgt	1800
ggaagcaaac	atcggcgggc	tacagtcagc	tcacagttca	aggttgactc	actgcattcc	1860
ttaatggcaa	cgctaagctc	ctctaatacct	ttctttgttc	gctgtatcaa	gccaaacatg	1920
cagaagatgc	cagaccagtt	tgaccaggcg	gttgtgctga	accagctgcg	gtactcaggg	1980
atgctggaga	ctgtgagaat	ccgcaaagct	gggtatgcgg	tccgaagacc	ctttcaggac	2040
ttttacaaaa	ggtataaagt	gctgatgagg	aatctggctc	tgcctgagga	cgtccgaggg	2100
aagtgcacga	gcctgctgca	gctctatgat	gcctccaaca	gcgagtggca	gctggggaag	2160
accaaggtat	ttcttcgaga	atccttggaa	cagaaactgg	agaagcggag	ggaagaggaa	2220
gtgagccacg	cggccatggt	gattcggggc	catgtcttgg	gcttcttagc	acggaaacaa	2280
tacagaaagg	tcctttattg	tgtggtgata	atacagaaga	attacagagc	attccttctg	2340
aggaggagat	ttttgcacct	gaaaaaggca	gccatagttt	tccagaagca	actcagaggt	2400
cagattgctc	ggagagttta	cagacaattg	ctggcagaga	aaagggagca	agaagaaaag	2460
aagaaacagg	aagaggaaga	aaagaagaaa	cgggaggaag	aagaaagaga	aagagagaga	2520
gagcgaagag	aagccgagct	ccgcgcccag	caggaagaag	aaacgaggaa	gcagcaagaa	2580
ctcgaagcct	tgcagaagag	ccagaaggaa	gctgaactga	cccgtgaact	ggagaaacag	2640
aaggaaaata	agcaggtgga	agagatcctc	cgtctggaga	aagaaatcga	ggacctgcag	2700

cgcatgaagg	agcagcagga	gctgtcgctg	accgaggctt	ccctgcagaa	gctgcaggag	2760
cggcgggacc	aggagctccg	caggctggag	gaggaagcgt	gcagggcggc	ccaggagttc	2820
ctcgagtccc	tcaatttcga	cgagatcgac	gagtgtgtcc	ggaatatcga	gcggctccctg	2880
tcggggggaa	gcgaattttc	cagcgagctg	gctgagagcg	catgcgagga	gaagcccaac	2940
ttcaacttca	gccagcccta	cccagaggag	gaggtcgatg	agggcttcga	agccgacgac	3000
gacgccttca	aggactcccc	caaccccgag	gagcacggcc	actcagacca	gcgaacaagt	3060
ggcatccgga	ccagcgatga	ctcttcagag	gaggacccat	acatgaacga	cacggtggtg	3120
cccaccagcc	ccagtgcgga	cagcacgggtg	ctgctcgccc	catcagtgcg	ggactccggg	3180
agcctacaca	actcctccag	cggcgagctc	acctactgca	tgccccagaa	cgctgggggac	3240
ttgccctccc	cagacggcga	ctacgactac	gaccaggatg	actatgagga	cggtgccatc	3300
acttcgggca	gcagcgtgac	cttctccaac	tcctacggca	gccagtggtc	ccccgactac	3360
cgctgctctg	tggggaccta	caacagctcg	ggtgcctacc	ggttcagctc	tgagggggcg	3420
cagtcctcgt	ttgaagatag	tgaagaggag	tttgattcca	ggtttgatac	agatgatgag	3480
ctttcatacc	ggcgtgactc	tgtgtacagc	tgtgtcactc	tgccgtatct	ccacagcttt	3540
ctgtacatga	aaggtggcct	gatgaactct	tggaaacgcc	gctgggtcgt	cctcaaggat	3600
gaaaccttct	tgtggttccg	ctccaagcag	gaggccctca	agcaaggctg	gctccacaaa	3660
aaaggggggg	gctcctccac	gctgtccagg	agaaattgga	agaagcgctg	gtttgtcctc	3720
cgccagtcca	agctgatgta	ctttgaaaac	gacagcgagg	agaagctcaa	gggcaccgta	3780
gaagtgcgaa	cggcaaaaaga	gatcatagat	aacaccacca	aggagaatgg	gatcgacatc	3840
attatggccg	ataggacttt	ccacctgatt	gcagagtccc	cagaagatgc	cagccagtgg	3900
ttcagcgtgc	tgagtcagggt	ccacgcgtcc	acggaccagg	agatccagga	gatgcatgat	3960
gagcaggcaa	acccacagaa	tgctgtgggc	accttgatg	tggggctgat	tgattctgtg	4020
tgtgcctcga	acagccctga	tagacccaac	tcgtttgtga	tcatcacggc	caaccgggtg	4080
ctgactgca	acgcgcacac	gccggaggag	atgcaccact	ggataaccct	gctgcagagg	4140
tccaaagggg	acaccagagt	ggagggccag	gaattcatcg	tgagaggatg	gttgacaaaa	4200
gaggtgaaga	acagtccaaa	gatgtcttca	ctgaaactga	agaaacgggtg	gtttgtactc	4260
accacacaatt	ccctggatta	ctacaagagt	tcagagaaga	acgcgctcaa	actggggacc	4320
ctggtcctca	acagcctctg	ctctgtcgtc	ccccagatg	agaagatatt	caaagagaca	4380
ggctactgga	acgtcacctg	gtacggggcg	aagcactggt	accggctcta	caccaagctg	4440
ctcaacgagg	ccaccgggtg	gtccagtgtc	attcaaaacg	tgactgacac	caaggccccg	4500
atcgacaccc	ccaccagca	gctgattcaa	gatatcaagg	agaactgcct	gaactcggat	4560
gtggtgggaa	agatttataa	gcggaacccg	atccttcgat	acacccatca	ccccttgac	4620
tccccgctcc	tgcccccttc	gtatggggac	ataaatctca	acttgctgaa	agacaaaagg	4680
tataccaccc	ttcaggatga	ggccatcaag	atattcaatt	ccctgcagca	actggagtcc	4740
atgtctgacc	caattccaat	aatccagggc	atcctacaga	cagggcatga	cctgcgacct	4800
ctgcgggacg	agctgtactg	ccagcttatc	aaacagacca	acaaagtgcc	ccaccccggc	4860
agtgtgggca	acctgtacag	ctggcagatc	ctgacatgcc	tgagctgcac	cttctctgcg	4920
agtcgagggg	ttctcaagta	tctcaagttc	catctgaaaa	ggatacggga	acagtttcca	4980
ggaaccgaga	tggaaaaata	cgctctcttc	acttaagaa	ctcttaagaa	aaccaaagtc	5040
cgagagtttg	tgccttcccg	agatgaaata	gaagctctga	tccacaggca	ggaaatgaca	5100
tccagcgtct	attgccatgg	cggcggctcc	tgcaagatca	ccatcaactc	ccacaccacc	5160
gctggggagg	tggtggagaa	gctgatccga	ggcctggcca	tggaggacag	caggaacatg	5220
tttgctttgt	ttgaatacaa	cggccacgtc	gacaaagcca	ttgaaagtcg	aaccgtcgta	5280
gctgatgtct	tagccaagtt	tgaaaaagctg	gctgccacat	ccgaggttgg	ggacctgcca	5340
tggaaattct	acttcaaact	ttactgcttc	ctggacacag	acaacgtgcc	aaaagacagt	5400
gtggagtttg	catttatggt	tgaacaggcc	cacgaagcgg	ttatccatgg	ccaccatcca	5460
gccccggaag	aaaacctcca	ggttcttgct	gcctgcgac	tccagtatct	gcagggggat	5520
tatactctgc	acgctgccat	cccacctctc	gaagagggtt	attccctgca	gagactcaag	5580
gcccgcacac	gccagtcaac	caaaaccttc	accccttggt	aacggctgga	gaagaggcgg	5640
acgagcttcc	tagaggggac	cctgaggcgg	agcttcggga	caggatccgt	ggtccggcag	5700
aaggtcgagg	aggagcagat	gctggacatg	tggattaagg	aagaagtctc	ctctgctcga	5760
gccagtatca	ttgacaagtg	gaggaaattt	cagggaatga	accaggaaca	ggccatggcc	5820
aagtacatgg	ccttgatcaa	ggagtggcct	ggctatggct	cgacgctggt	tgatgtggag	5880
tgcaagggaag	gtggcttccc	tcaggaactc	tggttgggtg	tcagcgcgga	cgccgtctcc	5940
gtctacaagc	gtggagaggg	aagaccactg	gaagtcttcc	agtatgaaca	cactctctct	6000
tttggggcac	ccctggcgaa	tacgtataag	atcgtggctg	atgagaggga	gctgctcttt	6060
gaaaccagtg	aggtagtgga	tgtggccaag	ctcatgaaag	cctacatcag	catgatcgtg	6120
aagaagcgct	acagcacgac	acgctccgcc	agcagccagg	gcagctccag	g	6171

<210> 10

<211> 2057

<212> PRT

<213> Homo sapiens

<400> 10

Phe	Cys	Leu	Gln	Gly	Thr	Arg	Val	Trp	Leu	Arg	Glu	Asn	Gly	Gln	His
1				5					10					15	

Phe	Pro	Ser	Thr	Val	Asn	Ser	Cys	Ala	Glu	Gly	Ile	Val	Val	Phe	Arg
			20					25					30		

Thr	Asp	Tyr	Gly	Gln	Val	Phe	Thr	Tyr	Lys	Gln	Ser	Thr	Ile	Thr	His
		35					40					45			

Gln	Lys	Val	Thr	Ala	Met	His	Pro	Thr	Asn	Glu	Glu	Gly	Val	Asp	Asp
	50					55					60				

Met	Ala	Ser	Leu	Thr	Glu	Leu	His	Gly	Gly	Ser	Ile	Met	Tyr	Asn	Leu
65					70					75					80

Phe	Gln	Arg	Tyr	Lys	Arg	Asn	Gln	Ile	Trp	Thr	Tyr	Ile	Gly	Ser	Ile
				85					90					95	

Leu	Ala	Ser	Val	Asn	Pro	Tyr	Gln	Pro	Ile	Ala	Gly	Leu	Tyr	Glu	Pro
			100					105					110		

Ala	Thr	Met	Glu	Gln	Tyr	Ser	Arg	Arg	His	Leu	Gly	Glu	Leu	Pro	Pro
		115					120					125			

His	Ile	Phe	Ala	Ile	Ala	Asn	Glu	Cys	Tyr	Arg	Cys	Leu	Trp	Lys	Arg
	130					135					140				

His	Asp	Asn	Gln	Cys	Ile	Leu	Ile	Lys	Gly	Glu	Ser	Gly	Ala	Gly	Lys
145					150					155					160

Thr	Glu	Ser	Thr	Lys	Leu	Ile	Leu	Lys	Phe	Leu	Ser	Val	Ile	Ser	Gln
				165					170					175	

Gln	Ser	Leu	Glu	Leu	Ser	Leu	Lys	Glu	Lys	Thr	Ser	Cys	Val	Glu	Arg
			180					185					190		

Ala	Ile	Leu	Glu	Ser	Ser	Pro	Ile	Met	Glu	Ala	Phe	Gly	Asn	Ala	Lys
		195					200					205			

Thr	Val	Tyr	Asn	Asn	Asn	Ser	Ser	Arg	Phe	Gly	Lys	Phe	Val	Gln	Leu
	210					215					220				

Asn	Ile	Cys	Gln	Lys	Gly	Asn	Ile	Gln	Gly	Gly	Arg	Ile	Val	Asp	Cys
225					230					235					240

Ile	Leu	Ser	Ser	Gln	Asn	Arg	Val	Val	Arg	Gln	Asn	Pro	Gly	Glu	Arg
				245					250					255	

Asn	Tyr	His	Ile	Phe	Tyr	Ala	Leu	Leu	Ala	Gly	Leu	Glu	His	Glu	Glu
			260					265					270		

Arg Glu Glu Phe Tyr Leu Ser Thr Pro Glu Asn Tyr His Tyr Leu Asn
 275 280 285
 Gln Ser Gly Cys Val Glu Asp Lys Thr Ile Ser Asp Gln Glu Ser Phe
 290 295 300
 Arg Glu Val Ile Thr Ala Met Asp Val Met Gln Phe Ser Lys Glu Glu
 305 310 315 320
 Val Arg Glu Val Ser Arg Leu Leu Ala Gly Ile Leu His Leu Gly Asn
 325 330 335
 Ile Glu Phe Ile Thr Ala Gly Gly Ala Gln Val Ser Phe Lys Thr Ala
 340 345 350
 Leu Gly Arg Ser Ala Glu Leu Leu Gly Leu Asp Pro Thr Gln Leu Thr
 355 360 365
 Asp Ala Leu Thr Gln Arg Ser Met Phe Leu Arg Gly Glu Glu Ile Leu
 370 375 380
 Thr Pro Leu Asn Val Gln Gln Ala Val Asp Ser Arg Asp Ser Leu Ala
 385 390 395 400
 Met Ala Leu Tyr Ala Cys Cys Phe Glu Trp Val Ile Lys Lys Ile Asn
 405 410 415
 Ser Arg Ile Lys Gly Asn Glu Asp Phe Lys Ser Ile Gly Ile Leu Asp
 420 425 430
 Ile Phe Gly Phe Glu Asn Phe Glu Val Asn His Phe Glu Gln Phe Asn
 435 440 445
 Ile Asn Tyr Ala Asn Glu Lys Leu Gln Glu Tyr Phe Asn Lys His Ile
 450 455 460
 Phe Ser Leu Glu Gln Leu Glu Tyr Ser Arg Glu Gly Leu Val Trp Glu
 465 470 475 480
 Asp Ile Asp Trp Ile Asp Asn Gly Glu Cys Leu Asp Leu Ile Glu Lys
 485 490 495
 Lys Leu Gly Leu Leu Ala Leu Ile Asn Glu Glu Ser His Phe Pro Gln
 500 505 510
 Ala Thr Asp Ser Thr Leu Leu Glu Lys Leu His Ser Gln His Ala Asn
 515 520 525
 Asn His Phe Tyr Val Lys Pro Arg Val Ala Val Asn Asn Phe Gly Val
 530 535 540
 Lys His Tyr Ala Gly Glu Val Gln Tyr Asp Val Arg Gly Ile Leu Glu
 545 550 555 560
 Lys Asn Arg Asp Thr Phe Arg Asp Asp Leu Leu Asn Leu Leu Arg Glu
 565 570 575

Ser	Arg	Phe	Asp	Phe	Ile	Tyr	Asp	Leu	Phe	Glu	His	Val	Ser	Ser	Arg	
			580					585					590			
Asn	Asn	Gln	Asp	Thr	Leu	Lys	Cys	Gly	Ser	Lys	His	Arg	Arg	Pro	Thr	
		595					600					605				
Val	Ser	Ser	Gln	Phe	Lys	Val	Asp	Ser	Leu	His	Ser	Leu	Met	Ala	Thr	
	610					615					620					
Leu	Ser	Ser	Ser	Asn	Pro	Phe	Phe	Val	Arg	Cys	Ile	Lys	Pro	Asn	Met	
625					630					635					640	
Gln	Lys	Met	Pro	Asp	Gln	Phe	Asp	Gln	Ala	Val	Val	Leu	Asn	Gln	Leu	
				645					650					655		
Arg	Tyr	Ser	Gly	Met	Leu	Glu	Thr	Val	Arg	Ile	Arg	Lys	Ala	Gly	Tyr	
			660					665					670			
Ala	Val	Arg	Arg	Pro	Phe	Gln	Asp	Phe	Tyr	Lys	Arg	Tyr	Lys	Val	Leu	
		675					680					685				
Met	Arg	Asn	Leu	Ala	Leu	Pro	Glu	Asp	Val	Arg	Gly	Lys	Cys	Thr	Ser	
	690					695					700					
Leu	Leu	Gln	Leu	Tyr	Asp	Ala	Ser	Asn	Ser	Glu	Trp	Gln	Leu	Gly	Lys	
705					710					715					720	
Thr	Lys	Val	Phe	Leu	Arg	Glu	Ser	Leu	Glu	Gln	Lys	Leu	Glu	Lys	Arg	
				725					730					735		
Arg	Glu	Glu	Glu	Val	Ser	His	Ala	Ala	Met	Val	Ile	Arg	Ala	His	Val	
			740					745					750			
Leu	Gly	Phe	Leu	Ala	Arg	Lys	Gln	Tyr	Arg	Lys	Val	Leu	Tyr	Cys	Val	
		755					760					765				
Val	Ile	Ile	Gln	Lys	Asn	Tyr	Arg	Ala	Phe	Leu	Leu	Arg	Arg	Arg	Phe	
	770					775						780				
Leu	His	Leu	Lys	Lys	Ala	Ala	Ile	Val	Phe	Gln	Lys	Gln	Leu	Arg	Gly	
785					790					795					800	
Gln	Ile	Ala	Arg	Arg	Val	Tyr	Arg	Gln	Leu	Leu	Ala	Glu	Lys	Arg	Glu	
				805					810					815		
Gln	Glu	Glu	Lys	Lys	Lys	Gln	Glu	Glu	Glu	Glu	Lys	Lys	Lys	Arg	Glu	
			820					825						830		
Glu	Glu	Glu	Arg	Glu	Arg	Glu	Arg	Glu	Arg	Arg	Glu	Ala	Glu	Leu	Arg	
		835					840					845				
Ala	Gln	Gln	Glu	Glu	Glu	Thr	Arg	Lys	Gln	Gln	Glu	Leu	Glu	Ala	Leu	
						855					860					
Gln	Lys	Ser	Gln	Lys	Glu	Ala	Glu	Leu	Thr	Arg	Glu	Leu	Glu	Lys	Gln	
865					870					875					880	

Lys	Glu	Asn	Lys	Gln	Val	Glu	Glu	Ile	Leu	Arg	Leu	Glu	Lys	Glu	Ile	
				885					890					895		
Glu	Asp	Leu	Gln	Arg	Met	Lys	Glu	Gln	Gln	Glu	Leu	Ser	Leu	Thr	Glu	
				900					905					910		
Ala	Ser	Leu	Gln	Lys	Leu	Gln	Glu	Arg	Arg	Asp	Gln	Glu	Leu	Arg	Arg	
				915					920					925		
Leu	Glu	Glu	Glu	Ala	Cys	Arg	Ala	Ala	Gln	Glu	Phe	Leu	Glu	Ser	Leu	
				930					935					940		
Asn	Phe	Asp	Glu	Ile	Asp	Glu	Cys	Val	Arg	Asn	Ile	Glu	Arg	Ser	Leu	
				945					950					955		
Ser	Gly	Gly	Ser	Glu	Phe	Ser	Ser	Glu	Leu	Ala	Glu	Ser	Ala	Cys	Glu	
				965					970					975		
Glu	Lys	Pro	Asn	Phe	Asn	Phe	Ser	Gln	Pro	Tyr	Pro	Glu	Glu	Glu	Val	
				980					985					990		
Asp	Glu	Gly	Phe	Glu	Ala	Asp	Asp	Asp	Ala	Phe	Lys	Asp	Ser	Pro	Asn	
				995					1000					1005		
Pro	Ser	Glu	His	Gly	His	Ser	Asp	Gln	Arg	Thr	Ser	Gly	Ile	Arg	Thr	
				1010					1015					1020		
Ser	Asp	Asp	Ser	Ser	Glu	Glu	Asp	Pro	Tyr	Met	Asn	Asp	Thr	Val	Val	
				1025					1030					1035		
Pro	Thr	Ser	Pro	Ser	Ala	Asp	Ser	Thr	Val	Leu	Leu	Ala	Pro	Ser	Val	
				1045					1050					1055		
Gln	Asp	Ser	Gly	Ser	Leu	His	Asn	Ser	Ser	Ser	Gly	Glu	Ser	Thr	Tyr	
				1060					1065					1070		
Cys	Met	Pro	Gln	Asn	Ala	Gly	Asp	Leu	Pro	Ser	Pro	Asp	Gly	Asp	Tyr	
				1075					1080					1085		
Asp	Tyr	Asp	Gln	Asp	Asp	Tyr	Glu	Asp	Gly	Ala	Ile	Thr	Ser	Gly	Ser	
				1090					1095					1100		
Ser	Val	Thr	Phe	Ser	Asn	Ser	Tyr	Gly	Ser	Gln	Trp	Ser	Pro	Asp	Tyr	
				1105					1110					1115		
Arg	Cys	Ser	Val	Gly	Thr	Tyr	Asn	Ser	Ser	Gly	Ala	Tyr	Arg	Phe	Ser	
				1125					1130					1135		
Ser	Glu	Gly	Ala	Gln	Ser	Ser	Phe	Glu	Asp	Ser	Glu	Glu	Asp	Phe	Asp	
				1140					1145					1150		
Ser	Arg	Phe	Asp	Thr	Asp	Asp	Glu	Leu	Ser	Tyr	Arg	Arg	Asp	Ser	Val	
				1155					1160					1165		
Tyr	Ser	Cys	Val	Thr	Leu	Pro	Tyr	Phe	His	Ser	Phe	Leu	Tyr	Met	Lys	
				1170					1175					1180		

Gly Gly Leu Met Asn Ser Trp Lys Arg Arg Trp Cys Val Leu Lys Asp
 1185 1190 1195 1200
 Glu Thr Phe Leu Trp Phe Arg Ser Lys Gln Glu Ala Leu Lys Gln Gly
 1205 1210 1215
 Trp Leu His Lys Lys Gly Gly Gly Ser Ser Thr Leu Ser Arg Arg Asn
 1220 1225 1230
 Trp Lys Lys Arg Trp Phe Val Leu Arg Gln Ser Lys Leu Met Tyr Phe
 1235 1240 1245
 Glu Asn Asp Ser Glu Glu Lys Leu Lys Gly Thr Val Glu Val Arg Thr
 1250 1255 1260
 Ala Lys Glu Ile Ile Asp Asn Thr Thr Lys Glu Asn Gly Ile Asp Ile
 1265 1270 1275 1280
 Ile Met Ala Asp Arg Thr Phe His Leu Ile Ala Glu Ser Pro Glu Asp
 1285 1290 1295
 Ala Ser Gln Trp Phe Ser Val Leu Ser Gln Val His Ala Ser Thr Asp
 1300 1305 1310
 Gln Glu Ile Gln Glu Met His Asp Glu Gln Ala Asn Pro Gln Asn Ala
 1315 1320 1325
 Val Gly Thr Leu Asp Val Gly Leu Ile Asp Ser Val Cys Ala Ser Asp
 1330 1335 1340
 Ser Pro Asp Arg Pro Asn Ser Phe Val Ile Ile Thr Ala Asn Arg Val
 1345 1350 1355 1360
 Leu His Cys Asn Ala Asp Thr Pro Glu Glu Met His His Trp Ile Thr
 1365 1370 1375
 Leu Leu Gln Arg Ser Lys Gly Asp Thr Arg Val Glu Gly Gln Glu Phe
 1380 1385 1390
 Ile Val Arg Gly Trp Leu His Lys Glu Val Lys Asn Ser Pro Lys Met
 1395 1400 1405
 Ser Ser Leu Lys Leu Lys Lys Arg Trp Phe Val Leu Thr His Asn Ser
 1410 1415 1420
 Leu Asp Tyr Tyr Lys Ser Ser Glu Lys Asn Ala Leu Lys Leu Gly Thr
 1425 1430 1435 1440
 Leu Val Leu Asn Ser Leu Cys Ser Val Val Pro Pro Asp Glu Lys Ile
 1445 1450 1455
 Phe Lys Glu Thr Gly Tyr Trp Asn Val Thr Val Tyr Gly Arg Lys His
 1460 1465 1470
 Cys Tyr Arg Leu Tyr Thr Lys Leu Leu Asn Glu Ala Thr Arg Trp Ser
 1475 1480 1485

Ser Val Ile Gln Asn Val Thr Asp Thr Lys Ala Pro Ile Asp Thr Pro
 1490 1495 1500

Thr Gln Gln Leu Ile Gln Asp Ile Lys Glu Asn Cys Leu Asn Ser Asp
 1505 1510 1515 1520

Val Val Glu Gln Ile Tyr Lys Arg Asn Pro Ile Leu Arg Tyr Thr His
 1525 1530 1535

His Pro Leu His Ser Pro Leu Leu Pro Leu Pro Tyr Gly Asp Ile Asn
 1540 1545 1550

Leu Asn Leu Leu Lys Asp Lys Gly Tyr Thr Thr Leu Gln Asp Glu Ala
 1555 1560 1565

Ile Lys Ile Phe Asn Ser Leu Gln Gln Leu Glu Ser Met Ser Asp Pro
 1570 1575 1580

Ile Pro Ile Ile Gln Gly Ile Leu Gln Thr Gly His Asp Leu Arg Pro
 1585 1590 1595 1600

Leu Arg Asp Glu Leu Tyr Cys Gln Leu Ile Lys Gln Thr Asn Lys Val
 1605 1610 1615

Pro His Pro Gly Ser Val Gly Asn Leu Tyr Ser Trp Gln Ile Leu Thr
 1620 1625 1630

Cys Leu Ser Cys Thr Phe Leu Pro Ser Arg Gly Ile Leu Lys Tyr Leu
 1635 1640 1645

Lys Phe His Leu Lys Arg Ile Arg Glu Gln Phe Pro Gly Thr Glu Met
 1650 1655 1660

Glu Lys Tyr Ala Leu Phe Thr Tyr Glu Ser Leu Lys Lys Thr Lys Cys
 1665 1670 1675 1680

Arg Glu Phe Val Pro Ser Arg Asp Glu Ile Glu Ala Leu Ile His Arg
 1685 1690 1695

Gln Glu Met Thr Ser Thr Val Tyr Cys His Gly Gly Gly Ser Cys Lys
 1700 1705 1710

Ile Thr Ile Asn Ser His Thr Thr Ala Gly Glu Val Val Glu Lys Leu
 1715 1720 1725

Ile Arg Gly Leu Ala Met Glu Asp Ser Arg Asn Met Phe Ala Leu Phe
 1730 1735 1740

Glu Tyr Asn Gly His Val Asp Lys Ala Ile Glu Ser Arg Thr Val Val
 1745 1750 1755 1760

Ala Asp Val Leu Ala Lys Phe Glu Lys Leu Ala Ala Thr Ser Glu Val
 1765 1770 1775

Gly Asp Leu Pro Trp Lys Phe Tyr Phe Lys Leu Tyr Cys Phe Leu Asp
 1780 1785 1790

Thr Asp Asn Val Pro Lys Asp Ser Val Glu Phe Ala Phe Met Phe Glu
 1795 1800 1805
 Gln Ala His Glu Ala Val Ile His Gly His His Pro Ala Pro Glu Glu
 1810 1815 1820
 Asn Leu Gln Val Leu Ala Ala Leu Arg Leu Gln Tyr Leu Gln Gly Asp
 1825 1830 1835 1840
 Tyr Thr Leu His Ala Ala Ile Pro Pro Leu Glu Glu Val Tyr Ser Leu
 1845 1850 1855
 Gln Arg Leu Lys Ala Arg Ile Ser Gln Ser Thr Lys Thr Phe Thr Pro
 1860 1865 1870
 Cys Glu Arg Leu Glu Lys Arg Arg Thr Ser Phe Leu Glu Gly Thr Leu
 1875 1880 1885
 Arg Arg Ser Phe Arg Thr Gly Ser Val Val Arg Gln Lys Val Glu Glu
 1890 1895 1900
 Glu Gln Met Leu Asp Met Trp Ile Lys Glu Glu Val Ser Ser Ala Arg
 1905 1910 1915 1920
 Ala Ser Ile Ile Asp Lys Trp Arg Lys Phe Gln Gly Met Asn Gln Glu
 1925 1930 1935
 Gln Ala Met Ala Lys Tyr Met Ala Leu Ile Lys Glu Trp Pro Gly Tyr
 1940 1945 1950
 Gly Ser Thr Leu Phe Asp Val Glu Cys Lys Glu Gly Gly Phe Pro Gln
 1955 1960 1965
 Glu Leu Trp Leu Gly Val Ser Ala Asp Ala Val Ser Val Tyr Lys Arg
 1970 1975 1980
 Gly Glu Gly Arg Pro Leu Glu Val Phe Gln Tyr Glu His Ile Leu Ser
 1985 1990 1995 2000
 Phe Gly Ala Pro Leu Ala Asn Thr Tyr Lys Ile Val Val Asp Glu Arg
 2005 2010 2015
 Glu Leu Leu Phe Glu Thr Ser Glu Val Val Asp Val Ala Lys Leu Met
 2020 2025 2030
 Lys Ala Tyr Ile Ser Met Ile Val Lys Lys Arg Tyr Ser Thr Thr Arg
 2035 2040 2045
 Ser Ala Ser Ser Gln Gly Ser Ser Arg
 2050 2055

<210> 11

<211> 9621

<212> DNA

<213> Homo sapiens

<400> 11

```

agctagtatc ttttattgtc agaacttctg tgagccaaca aacagttttg catggttgta 60
cacaaagggg caaggcaaat ttcttttttc gtgtgggtag acttagttgg cccaagtcct 120
taaaactttt ccatataaaa ataaaaagtc caagaccaga ttatttttct tctggtcata 180
aatgctgatt tatttacagg tgccttggtc agaccaccat tataaacttg ggataaaaata 240
tgtgtgtatt aaagcctcag catttaatgt cagggtcctt tgaagattca ctcaagtgtt 300
aagacgtttc tggaatgcag cgtctctccc ccatagtcaa catggttatt atatctgtaa 360
tctatccaga atgatagaag ctaaccttcc aagtaacact ttgtttttta cttaaactctt 420
ttagacatga aagactccaa aatgacttca ttcttggtct aaaaccagca ctggagccag 480
ctgttgaaaga gtggtttata aatacagtta tcttgtaggc tgcttatctg tttataatac 540
agcagacaca gatggcagac tttgctacat gtaaaacaat ggagtcaaca cgtgtttttc 600
aaaatacagc aaagacagga aaatccagga tttggggttg ttaataaaac caccttataa 660
agtaacaatt gagactatag ctctgcatta ttaaaatata cagactgtgt acaccattac 720
acatcctttt tccctttgct ttttaatgct catgaaacca tgattaaggt gttgagttta 780
tgaacacatg cacgaacagg caagcacgta cacttaaaag atgaaacaaa gaaaaagtt 840
gattcatgtc attccatgag aaaggctgcc cgcagcactc cagctcaaac acactgtccc 900
ctcagactct ccatccccct tcccactccc tcaccttccc tcagattcgg ggaaatcagg 960
ttgggaggtt agtgcacat tgacagagaa tgccccctt ccacgctctg ttaagtctcc 1020
cccagaaggg ggaaaggcag ttcccttcag tagcacagtt acggtcgatt agtgttggtt 1080
ccacaagtta aggcacttcc ggctgctttg gtggcagcgt gggtccctccc ctcccttttt 1140
aaggcatgtg tcctctaaga gtagtaaagc tttggaaact gtgcagactg ttaaagttga 1200
cagcttaata caggatcaat gaaggcggga ggcaaaagga tcctcggaga cactccctc 1260
agaccagaag ctccagaaa gcctgggcag ctctgtgttt gttttggctg ggcatggcac 1320
actggagcca gcctaggcca gaggggtggtg cggtcaggta gcaaagacag gtgggctctg 1380
tcccgcttc acctggagct gccctggctg ctggcggagc gtgtcgtgct gtagcgcttc 1440
ttcacgatca tgetgatgta ggctttcatg agcttgacca catccaccac ctactgggtt 1500
tcaaagagca gtcctctctc atcgaccacg atcttatacg tattcgccag ggttgcccca 1560
aaagagagga tgtgttcata ctggaagact tccagtggtc tccctctccc acgcttgtag 1620
acggagacgg cgtccgcgct gacacccaac cagagttcct gaggggaagcc accttctctg 1680
cactgggtgg caagagtcaa cagaagagtt aagtcatgaa gtggttgga acagaaagca 1740
tctaaaccat aagacaggct ttgagtgaag tcctctgtgc agaagattaa atatattcga 1800
tgtgcatgca tgcattggag ggctgaaat atgaaaaatg gcacctctct ggctatcttg 1860
atttctaact agttaatctc acgcttttgg gaaaacctca ctaactggca gactctaaca 1920
tcttgctttg actctccact tctcagcatt attctactag ctgtttggat tagctacgtg 1980
gaagtggcct ggaaacgtac atgcttgccc gggggactta agaaagcttc cctgcaaccc 2040
aagccaagtc tactcttgta ttaatatctc cagttctgcc tccaatcctc tttgcggatg 2100
gttagtcttc aaatacaaaa tctaggatca cagaggaaaa ttctccaaat cagcatgct 2160
gagcagttct ggctcctctt cacaaggagc agcaatggcc ttccatagc agagtgggaa 2220
cagggacttt accagtttaa ctgtagactt tctgtacag attggtggaa gaaaataaga 2280
ccccatatga aggggctaac aacacagggc tgatccaaac ctggacaagc aggagggcta 2340
taaatggag acgtgaaaa gagtctctag tttatatccc taataaccag acattctctg 2400
catcctccat gcaaaagcca gtagctttct ttttttcttt ttttttttga cggagtctca 2460
ctctgtcgcc caggctggag tgcagtggcg tgatcctggc tcaactgcaac ctccacctcc 2520
tgagttcaag cgattctcct gtctcagcct ctcgagtagc tgggattaca ggtgcatgcc 2580
accacgcccc gctaattttt tgcagagatg gggtttcacc gtgttagcca ggatggtctc 2640
gatctcctga cctcatgatc cgcccgccct ggccctccaa agcgctggga ttacaggcat 2700
gagccaccgc gcccgcccaa gccagtagct ttctatgcta attcacagct caggttttgc 2760
aggaagccaa gattttaact gctattatct attccttgct agggagaaat ggaattatgg 2820
ctttgtacaa agcacctgat ttttttatac ttaaaaacag gcataattga accaaccaaa 2880
ccaatcaaaa acatcaccta atgaaaagcc acccaggat tctagaattt ataattttta 2940
gaattttata cagcctcaat ataaagtcac cagatatagc ctgaattact gtgatcataa 3000
aaaatggaag ctaatctaga cgatgagctg gcacacttat ctgttaaggg ctgcatagta 3060
aagattttta gactttgtgg atcacatggt ctctgtcaca actactcaac tctgtacaaa 3120
aacagccagg gaatatatct aaaggaatga gcttggtgtg gttccaataa aactttgttt 3180
agaaaaaag gaggcaggca agatctgacc cacagaccag tttgccaac tctcatctag 3240
acaattagta agatttcttt tcaataagcg gtctacttaa aacaaaacaa aaatcagtac 3300
tgggttgatg ccaatggcta aattccatta cgagatagac attcttctt tcaaacagat 3360
ggctgtaaag aaaaaacaaa gtaaaatgca agtatatcca aagtttctaa tttgtatata 3420

```

cagctataac	atTTTTTTaa	atgtagattt	ttatcagtg	ttaaaaaatt	agatctatag	3480
cttccctaag	gaagggtaga	agaatagatg	acatcttaat	tttgcattca	ttcctaatat	3540
tacagatgca	tttactacac	aggagaagag	aaactgtgag	gagaagggag	gcgttaatgg	3600
tacaattttg	ggggctcgaa	aaaaagaggt	tgagagagca	aaatgctcca	tcttgtcttc	3660
tctccacatg	aacttggccg	tgatccatgt	tctcagatgc	cagcaccag	cccaccccaa	3720
cacatggcag	ccagttctca	cctccacatc	aaacagcgtc	gagccatagc	caggccactc	3780
cttgatcaag	gccatgtact	tggccatggc	ctgttctctg	ttcattccct	gaaatttctc	3840
ccacttgtca	atgatactgg	ctcgagcaga	ggagacttct	tccttaatcc	acatgtccag	3900
catctgtctc	tcctcgacct	tctgccggac	cacggatcct	gtccggaagc	tccgcctcag	3960
ggtccccctc	aggaagctcg	tccgcctctt	ctccagccgt	tcacaagggg	tgaaggtttt	4020
ggttgactgg	ctgatgcggg	ccttgagctc	ctgcagggaa	taaacctctt	cgagaggtgg	4080
gatggcagcg	tgacagatat	aatccccctg	cagatactgg	agtcgcaggg	cagcaagaac	4140
ctggaggttt	tcttccgggg	ctggatgggt	gccatggata	accgcttcgt	gggcctgttc	4200
aaacataaat	gcaaaactcca	cactgtcttt	tggcacggtg	tctgtgtcca	ggaagcagta	4260
aagtttgaag	tagaatttcc	atggcaggtc	cccaacctcg	gatgtggcag	ccagcttttc	4320
aaacttggct	aagacatcag	ctacgacggg	tcgactttca	atggctttgt	cgacgtggcc	4380
gttgatttca	aacaaagcaa	acatgttctc	gctgtcctcc	atggccaggc	ctcggatcag	4440
cttctccacc	acctccccag	cgggtggtgtg	ggagttgatg	gtgatcttgc	aggagccgcc	4500
gccatggcaa	tagaccgtgg	atgtcatattc	ctgcctgtgg	atcagagctt	ctatttcatc	4560
tcgggaaggc	acaaactctc	ggcatttggg	tttcttaaga	gattcgtaag	tgaagagagc	4620
gtatttttcc	atctcggttc	ctggaaactg	ttcccgtatc	cttttcagat	ggaacttgag	4680
atacttgaga	atcccccgac	tcggcaggaa	ggtgcagctc	aggcatgtca	ggatctgcca	4740
gctgtacagg	ttgccacac	tgccgggggtg	gggcactttg	ttggtctgtt	tgataagctg	4800
gcagtacagc	tcgtcccgc	gaggtcgcag	gtcatgccct	gtctgtagga	tgccctggat	4860
tattggaatt	gggtcagaca	tggactccag	ttgctgcagg	gaattgaata	tcttgatggc	4920
ctcatcctga	aggggtggtat	agcctttgtc	tttcagcaag	ttgagattta	tgtccccata	4980
cgggaagggc	aggagcgggg	agtgcgaagg	gtgatgggtg	tatcgaagga	tcgggttccg	5040
cttgtaaatac	tggtccacca	catccgagtt	caggcagttc	tccttgatat	cttgaatcag	5100
ctgctgggtg	gggggtgtcg	tcggggcctt	ggtgtcagtc	acgttttgaa	tgacactgga	5160
ccaccgggtg	gcctcgttga	gcagcttggt	gtagagccgg	taacagtgtc	tcgcccgtga	5220
cacggtgacg	ttccagtagc	ctgtctcttt	gaatatcttc	tcactctggg	ggacgacaga	5280
gcagaggctg	ttgaggacca	gggtccccag	tttgagcgcg	ttcttctctg	aactcttgta	5340
gtaatccagg	gaattgtggg	tgagtacaaa	ccaccgtttc	ttcagtttca	gtgaagacat	5400
ctttggactg	ttcttcacct	ctttgtgcaa	ccatcctctc	acgatgaatt	cctggccctc	5460
cactctgggtg	ttccccttgg	acctctgcag	cagggttatc	cagtgggtga	tctcctccgg	5520
cgtgtcggcg	ttgcagtgca	gcacccgggt	ggcgtgatg	atcacaaacg	agttgggtct	5580
atcagggctg	tcagaggcac	acacagaatc	aatcagcccc	acatccaagg	tgcccacagc	5640
attctgtggg	tttgctgtct	catcatgcat	ctcctggatc	tcctgggtccg	tggacgcgtg	5700
gacctgactc	agcacgttga	accactgggt	ggcatcttct	ggggactctg	caatcagggtg	5760
gaaagtcccta	tcggccataa	tgatgtcgat	cccattctcc	ttggtgggtg	tatctatgat	5820
ctcttttgcc	gttcgcactt	ctacgggtgcc	cttgagcttc	tcctcgtgtg	cgttttcaaa	5880
gtacatcagc	ttggactggc	ggaggacaaa	ccagcgttcc	ttccaatttc	tcctggacag	5940
cgtggaggag	cccccccctt	ttttgtggag	ccagccttgc	ttgaggccct	cctgcttggg	6000
gcggaaccac	aagaaggttt	catccttgag	gacgcaccag	cggcgtttcc	aagagttcat	6060
caggccacct	ttcatgtaca	gaaagctgtg	gaaatacggc	agagtgcac	agctgtacac	6120
agagtcacgc	cggtatgaaa	gctcatcatc	tgtatcaaac	ctggaatcaa	agtcctcttc	6180
actatcttca	aacgaggact	gcgccccctc	agagctgaac	cggtaggcac	ccgagctgtt	6240
gtaggtcccc	acagagcagc	ggtagtcggg	ggacctagg	ctgccgtagg	agttggagaa	6300
ggtcacgctg	ctgccggaag	tgatggcacc	gtcctcatag	tcactctggg	cgtagtcgta	6360
gtcgcctgtc	ggggagggca	agtccccagc	gttctggggc	atgcagtagg	tggactcgcc	6420
gctggaggag	ttgtgtaggc	tcccggagtc	ctgcactgat	ggggcgagca	gcaccgtgct	6480
gtccgcactg	gggctgggtg	gcaccaccgt	gtcgttcatg	tatgggtcct	cctctgaaga	6540
gtcatcgctg	gtccggatgc	cacttggttcg	ctggtctgag	tggcgtgct	cgctgggggt	6600
gggggagtc	ttgaaggcgt	cgtcgtcggc	ttcgaagccc	tcactgcacct	cctcctctgg	6660
gtagggctgg	ctgaagttga	agttgggctt	ctcctcgcat	gcgtctcag	ccagctcgct	6720
ggaaaattcg	cttccccccg	acagggaccg	ctcgatatcc	cggacacact	cgtcgatctc	6780
gtcgaaattg	agggactcga	ggaactcctg	ggcgcctctg	cacgttctct	cctccagcct	6840
gcggagctcc	tggtcccgc	gctcctgcag	cttctgcagg	gaagcctcgg	tcagcgacag	6900

```

ctcctgctgc tccttcatgc gctgcaggtc ctcgatttct ttctccagac ggaggatctc 6960
ttccacctgc ttattttcct tctgtttctc cagttcacgg gtcagttcag ctcccttctg 7020
gctcttctgc aaggcttcga gttcttgctg ctccctcgtt tcttcttctt gctgggcgcg 7080
gagctcggct tctcttcgct ctctctctct ttctcttctt tcttctctcc gtttcttctt 7140
ttcttctctt tctgtttctt tcttttcttc ttgtccctt ttctctgcca gcaattgtct 7200
gtaaactctc cgagcaatct gacctctgag ttgtctctgg aaaactatgg ctgccttttt 7260
caggtgcaaa aatctcctcc tcagaaggaa tgctctgtaa ttcttctgta ttatcaccac 7320
acaataaagg acctttctgt attgtttccg tgctaagaag cccaagacat gggcccgaa 7380
caccatggcc gcgtggctca ctctctcttc cctccgcttc tccagtttct gttccaagga 7440
ttctcgaaga aataccttgg tcttccccag ctgccactcg ctggtggagg catcatagag 7500
ctgcagcagg ctcgctgact tccctcggac gtccctcaggc agagccagat tctcatcag 7560
cactttatac cttttgtaaa agtccgaaa gggctctcgg accgcatacc cagctttgcg 7620
gattctcaca gtctccagca tccctgagta ccgcagctgg ttcagcacia ccgcctggtc 7680
aaactggctt ggcattcttct gcatgtttgg cttgatacag cgaacaaaga aaggattaga 7740
ggagcttagc gttgccatta aggaatgcag tgagtcaacc ttgaactgtg agctgactgt 7800
aggccgcgca tgtttgcttc cacatttcaa ggtatcctgg ttggtgcggc ttgaaacatg 7860
ttcaaaaaga tcgtagataa agtcaaaccg gctttctctt agcaaattga gaaggtcatc 7920
tcgaaatgta tctctgttct tctccaagat acctcggaca tcatattgca cctctccagc 7980
atagtgtctc actccaaaat tgttaactgc aactctgggc ttacataaaa agtggttatt 8040
cgcattgtga ctgtgtagct tctccaataa ggtgctgtct gtggcttgag gaaaatggct 8100
ttcttcattg ataagggcta ggaggccaag tttcttctca atcaagtcca ggcattcttc 8160
attgtctatc cagtcaatat ctccccacac taatccttcc ctgctatatt ctagtgttcc 8220
taaagaaaaa atatgcttgt tgaagtactc ctgaagtttc tcgtttgcat agtttatatt 8280
gaactgttca aagtgtattaa cctcaaagtt ttcaaattcca aagatgtcga ggatgccaat 8340
agacttgaag tctcatttgc ctttgatcct gctgttgatc ttcttgatta cccactcaaa 8400
gcagcacgca tacagagcca tggccaggga gtccctgctg tctactgcct gttgaacatt 8460
gagaggcgtg aggatctctt ctccctgag gaacattgat ctctgggtca aagcatctgt 8520
gagctgtgtt ggggtccagcc caagtaactc cgcagatctg cccaaagctg ttttgaagga 8580
aacctgtgcc ccaccagcag tgataaaattc tatgttccca agatgcagta taccagcaag 8640
cagcctcgac acttcccga cttcctcctt gctgaactgc atcacgtcca ttgccgtaat 8700
aacttcccta aaggattcct ggtcactgat tgtcttgtct tctacacatc cagactgatt 8760
caagtagtgg tagttttctg gcgtagataa ataaaaattt tctctttctt catgttccag 8820
ccctgccagc agtgcataaa atatgtgata attcctttcc ccgggatttt gccttactac 8880
tcggttctgg gaagagagga tacaatctac aattctcccg ccctgaatat ttcctttctg 8940
acagatgttc agctgaacaa acttcccaaa gcgactagag ttgttggtgt acacggtctt 9000
cgcattgccg aaagcttcca tgatggggct gctttcaaga atagctcgtt caacacagga 9060
tgtcttctcc tttaaggaca attccaaaga ctggtgactg atgactgaca gaaacttgag 9120
gatcaattta gtgctttcgg ttttacctgc cccactttca cccttgatga ggatgcactg 9180
ggtgtcgtgg cgcttcacaa ggcagcggta gcactcgttg gcgatggcga agatgtgcgg 9240
gggcagctcg ccaggtggc gccggctgta ctgctccatg gtggcaggct cgtacagccc 9300
ggcgatgggc tggtaggggt tcacagaggc caggatggag ccgatgtagg tccatatttg 9360
atttctctta taccgctgga ataagttata catgatggag ccgccatgga gctctgtcaa 9420
ggacgccatg tcatccacgc cctcctcgtt cgtgggggtgc atagcagtca ccttctggtg 9480
ggtaattgtg ctctgcttgt aagtgaatac ctgaccatag tctgtccgga agacgacgat 9540
gccttctgca caggaattta cagtacttgg aaaatgctgg ccattttctc tcagccagac 9600
ccgtgttccc tgtaaacaaa a 9621

```

<210> 12

<211> 2048

<212> PRT

<213> Homo sapiens

<400> 12

Phe Cys Leu Gln Gly Thr Arg Val Trp Leu Arg Glu Asn Gly Gln His

1

5

10

15

Phe Pro Ser Thr Val Asn Ser Cys Ala Glu Gly Ile Val Val Phe Arg
 20 25 30
 Thr Asp Tyr Gly Gln Val Phe Thr Tyr Lys Gln Ser Thr Ile Thr His
 35 40 45
 Gln Lys Val Thr Ala Met His Pro Thr Asn Glu Glu Gly Val Asp Asp
 50 55 60
 Met Ala Ser Leu Thr Glu Leu His Gly Gly Ser Ile Met Tyr Asn Leu
 65 70 75 80
 Phe Gln Arg Tyr Lys Arg Asn Gln Ile Trp Thr Tyr Ile Gly Ser Ile
 85 90 95
 Leu Ala Ser Val Asn Pro Tyr Gln Pro Ile Ala Gly Leu Tyr Glu Pro
 100 105 110
 Ala Thr Met Glu Gln Tyr Ser Arg Arg His Leu Gly Glu Leu Pro Pro
 115 120 125
 His Ile Phe Ala Ile Ala Asn Glu Cys Tyr Arg Cys Leu Trp Lys Arg
 130 135 140
 His Asp Asn Gln Cys Ile Leu Ile Lys Gly Glu Ser Gly Ala Gly Lys
 145 150 155 160
 Thr Glu Ser Thr Lys Leu Ile Leu Lys Phe Leu Ser Val Ile Ser Gln
 165 170 175
 Gln Ser Leu Glu Leu Ser Leu Lys Glu Lys Thr Ser Cys Val Glu Arg
 180 185 190
 Ala Ile Leu Glu Ser Ser Pro Ile Met Glu Ala Phe Gly Asn Ala Lys
 195 200 205
 Thr Val Tyr Asn Asn Asn Ser Ser Arg Phe Gly Lys Phe Val Gln Leu
 210 215 220
 Asn Ile Cys Gln Lys Gly Asn Ile Gln Gly Gly Arg Ile Val Asp Cys
 225 230 235 240
 Ile Leu Ser Ser Gln Asn Arg Val Val Arg Gln Asn Pro Gly Glu Arg
 245 250 255
 Asn Tyr His Ile Phe Tyr Ala Leu Leu Ala Gly Leu Glu His Glu Glu
 260 265 270
 Arg Glu Glu Phe Tyr Leu Ser Thr Pro Glu Asn Tyr His Tyr Leu Asn
 275 280 285
 Gln Ser Gly Cys Val Glu Asp Lys Thr Ile Ser Asp Gln Glu Ser Phe
 290 295 300
 Arg Glu Val Ile Thr Ala Met Asp Val Met Gln Phe Ser Lys Glu Glu
 305 310 315 320

Val Arg Glu Val Ser Arg Leu Leu Ala Gly Ile Leu His Leu Gly Asn
 325 330 335
 Ile Glu Phe Ile Thr Ala Gly Gly Ala Gln Val Ser Phe Lys Thr Ala
 340 345 350
 Leu Gly Arg Ser Ala Glu Leu Leu Gly Leu Asp Pro Thr Gln Leu Thr
 355 360 365
 Asp Ala Leu Thr Gln Arg Ser Met Phe Leu Arg Gly Glu Glu Ile Leu
 370 375 380
 Thr Pro Leu Asn Val Gln Gln Ala Val Asp Ser Arg Asp Ser Leu Ala
 385 390 395 400
 Met Ala Leu Tyr Ala Cys Cys Phe Glu Trp Val Ile Lys Lys Ile Asn
 405 410 415
 Ser Arg Ile Lys Gly Asn Glu Asp Phe Lys Ser Ile Gly Ile Leu Asp
 420 425 430
 Ile Phe Gly Phe Glu Asn Phe Glu Val Asn His Phe Glu Gln Phe Asn
 435 440 445
 Ile Asn Tyr Ala Asn Glu Lys Leu Gln Glu Tyr Phe Asn Lys His Ile
 450 455 460
 Phe Ser Leu Glu Gln Leu Glu Tyr Ser Arg Glu Gly Leu Val Trp Glu
 465 470 475 480
 Asp Ile Asp Trp Ile Asp Asn Gly Glu Cys Leu Asp Leu Ile Glu Lys
 485 490 495
 Lys Leu Gly Leu Leu Ala Leu Ile Asn Glu Glu Ser His Phe Pro Gln
 500 505 510
 Ala Thr Asp Ser Thr Leu Leu Glu Lys Leu His Ser Gln His Ala Asn
 515 520 525
 Asn His Phe Tyr Val Lys Pro Arg Val Ala Val Asn Asn Phe Gly Val
 530 535 540
 Lys His Tyr Ala Gly Glu Val Gln Tyr Asp Val Arg Gly Ile Leu Glu
 545 550 555 560
 Lys Asn Arg Asp Thr Phe Arg Asp Asp Leu Leu Asn Leu Leu Arg Glu
 565 570 575
 Ser Arg Phe Asp Phe Ile Tyr Asp Leu Phe Glu His Val Ser Ser Arg
 580 585 590
 Asn Asn Gln Asp Thr Leu Lys Cys Gly Ser Lys His Arg Arg Pro Thr
 595 600 605
 Val Ser Ser Gln Phe Lys Val Asp Ser Leu His Ser Leu Met Ala Thr
 610 615 620

Leu Ser Ser Ser Asn Pro Phe Phe Val Arg Cys Ile Lys Pro Asn Met
 625 630 635 640
 Gln Lys Met Pro Asp Gln Phe Asp Gln Ala Val Val Leu Asn Gln Leu
 645 650 655
 Arg Tyr Ser Gly Met Leu Glu Thr Val Arg Ile Arg Lys Ala Gly Tyr
 660 665 670
 Ala Val Arg Arg Pro Phe Gln Asp Phe Tyr Lys Arg Tyr Lys Val Leu
 675 680 685
 Met Arg Asn Leu Ala Leu Pro Glu Asp Val Arg Gly Lys Cys Thr Ser
 690 695 700
 Leu Leu Gln Leu Tyr Asp Ala Ser Asn Ser Glu Trp Gln Leu Gly Lys
 705 710 715 720
 Thr Lys Val Phe Leu Arg Glu Ser Leu Glu Gln Lys Leu Glu Lys Arg
 725 730 735
 Arg Glu Glu Glu Val Ser His Ala Ala Met Val Ile Arg Ala His Val
 740 745 750
 Leu Gly Phe Leu Ala Arg Lys Gln Tyr Arg Lys Val Leu Tyr Cys Val
 755 760 765
 Val Ile Ile Gln Lys Asn Tyr Arg Ala Phe Leu Leu Arg Arg Arg Phe
 770 775 780
 Leu His Leu Lys Lys Ala Ala Ile Val Phe Gln Lys Gln Leu Arg Gly
 785 790 795 800
 Gln Ile Ala Arg Arg Val Tyr Arg Gln Leu Leu Ala Glu Lys Arg Glu
 805 810 815
 Gln Glu Glu Lys Lys Lys Gln Glu Glu Glu Lys Lys Lys Arg Glu
 820 825 830
 Glu Glu Glu Arg Glu Arg Glu Arg Glu Arg Arg Glu Ala Glu Leu Arg
 835 840 845
 Ala Gln Gln Glu Glu Glu Thr Arg Lys Gln Gln Glu Leu Glu Ala Leu
 850 855 860
 Gln Lys Ser Gln Lys Glu Ala Glu Leu Thr Arg Glu Leu Glu Lys Gln
 865 870 875 880
 Lys Glu Asn Lys Gln Val Glu Glu Ile Leu Arg Leu Glu Lys Glu Ile
 885 890 895
 Glu Asp Leu Gln Arg Met Lys Glu Gln Gln Glu Leu Ser Leu Thr Glu
 900 905 910
 Ala Ser Leu Gln Lys Leu Gln Glu Arg Arg Asp Gln Glu Leu Arg Arg
 915 920 925

Leu Glu Glu Glu Ala Cys Arg Ala Ala Gln Glu Phe Leu Glu Ser Leu
 930 935 940
 Asn Phe Asp Glu Ile Asp Glu Cys Val Arg Asn Ile Glu Arg Ser Leu
 945 950 955 960
 Ser Gly Gly Ser Glu Phe Ser Ser Glu Leu Ala Glu Ser Ala Cys Glu
 965 970 975
 Glu Lys Pro Asn Phe Asn Phe Ser Gln Pro Tyr Pro Glu Glu Glu Val
 980 985 990
 Asp Glu Gly Phe Glu Ala Asp Asp Asp Ala Phe Lys Asp Ser Pro Asn
 995 1000 1005
 Pro Ser Glu His Gly His Ser Asp Gln Arg Thr Ser Gly Ile Arg Thr
 1010 1015 1020
 Ser Asp Asp Ser Ser Glu Glu Asp Pro Tyr Met Asn Asp Thr Val Val
 1025 1030 1035 1040
 Pro Thr Ser Pro Ser Ala Asp Ser Thr Val Leu Leu Ala Pro Ser Val
 1045 1050 1055
 Gln Asp Ser Gly Ser Leu His Asn Ser Ser Ser Gly Glu Ser Thr Tyr
 1060 1065 1070
 Cys Met Pro Gln Asn Ala Gly Asp Leu Pro Ser Pro Asp Gly Asp Tyr
 1075 1080 1085
 Asp Tyr Asp Gln Asp Asp Tyr Glu Asp Gly Ala Ile Thr Ser Gly Ser
 1090 1095 1100
 Ser Val Thr Phe Ser Asn Ser Tyr Gly Ser Gln Trp Ser Pro Asp Tyr
 1105 1110 1115 1120
 Arg Cys Ser Val Gly Thr Tyr Asn Ser Ser Gly Ala Tyr Arg Phe Ser
 1125 1130 1135
 Ser Glu Gly Ala Gln Ser Ser Phe Glu Asp Ser Glu Glu Asp Phe Asp
 1140 1145 1150
 Ser Arg Phe Asp Thr Asp Asp Glu Leu Ser Tyr Arg Arg Asp Ser Val
 1155 1160 1165
 Tyr Ser Cys Val Thr Leu Pro Tyr Phe His Ser Phe Leu Tyr Met Lys
 1170 1175 1180
 Gly Gly Leu Met Asn Ser Trp Lys Arg Arg Trp Cys Val Leu Lys Asp
 1185 1190 1195 1200
 Glu Thr Phe Leu Trp Phe Arg Ser Lys Gln Glu Ala Leu Lys Gln Gly
 1205 1210 1215
 Trp Leu His Lys Lys Gly Gly Gly Ser Ser Thr Leu Ser Arg Arg Asn
 1220 1225 1230

Trp Lys Lys Arg Trp Phe Val Leu Arg Gln Ser Lys Leu Met Tyr Phe
 1235 1240 1245
 Glu Asn Asp Ser Glu Glu Lys Leu Lys Gly Thr Val Glu Val Arg Thr
 1250 1255 1260
 Ala Lys Glu Ile Ile Asp Asn Thr Thr Lys Glu Asn Gly Ile Asp Ile
 1265 1270 1275 1280
 Ile Met Ala Asp Arg Thr Phe His Leu Ile Ala Glu Ser Pro Glu Asp
 1285 1290 1295
 Ala Ser Gln Trp Phe Ser Val Leu Ser Gln Val His Ala Ser Thr Asp
 1300 1305 1310
 Gln Glu Ile Gln Glu Met His Asp Glu Gln Ala Asn Pro Gln Asn Ala
 1315 1320 1325
 Val Gly Thr Leu Asp Val Gly Leu Ile Asp Ser Val Cys Ala Ser Asp
 1330 1335 1340
 Ser Pro Asp Arg Pro Asn Ser Phe Val Ile Ile Thr Ala Asn Arg Val
 1345 1350 1355 1360
 Leu His Cys Asn Ala Asp Thr Pro Glu Glu Met His His Trp Ile Thr
 1365 1370 1375
 Leu Leu Gln Arg Ser Lys Gly Asp Thr Arg Val Glu Gly Gln Glu Phe
 1380 1385 1390
 Ile Val Arg Gly Trp Leu His Lys Glu Val Lys Asn Ser Pro Lys Met
 1395 1400 1405
 Ser Ser Leu Lys Leu Lys Lys Arg Trp Phe Val Leu Thr His Asn Ser
 1410 1415 1420
 Leu Asp Tyr Tyr Lys Ser Ser Glu Lys Asn Ala Leu Lys Leu Gly Thr
 1425 1430 1435 1440
 Leu Val Leu Asn Ser Leu Cys Ser Val Val Pro Pro Asp Glu Lys Ile
 1445 1450 1455
 Phe Lys Glu Thr Gly Tyr Trp Asn Val Thr Val Tyr Gly Arg Lys His
 1460 1465 1470
 Cys Tyr Arg Leu Tyr Thr Lys Leu Leu Asn Glu Ala Thr Arg Trp Ser
 1475 1480 1485
 Ser Val Ile Gln Asn Val Thr Asp Thr Lys Ala Pro Ile Asp Thr Pro
 1490 1495 1500
 Thr Gln Gln Leu Ile Gln Asp Ile Lys Glu Asn Cys Leu Asn Ser Asp
 1505 1510 1515 1520
 Val Val Glu Gln Ile Tyr Lys Arg Asn Pro Ile Leu Arg Tyr Thr His
 1525 1530 1535

His Pro Leu His Ser Pro Leu Leu Pro Leu Pro Tyr Gly Asp Ile Asn
 1540 1545 1550
 Leu Asn Leu Leu Lys Asp Lys Gly Tyr Thr Thr Leu Gln Asp Glu Ala
 1555 1560 1565
 Ile Lys Ile Phe Asn Ser Leu Gln Gln Leu Glu Ser Met Ser Asp Pro
 1570 1575 1580
 Ile Pro Ile Ile Gln Gly Ile Leu Gln Thr Gly His Asp Leu Arg Pro
 1585 1590 1595 1600
 Leu Arg Asp Glu Leu Tyr Cys Gln Leu Ile Lys Gln Thr Asn Lys Val
 1605 1610 1615
 Pro His Pro Gly Ser Val Gly Asn Leu Tyr Ser Trp Gln Ile Leu Thr
 1620 1625 1630
 Cys Leu Ser Cys Thr Phe Leu Pro Ser Arg Gly Ile Leu Lys Tyr Leu
 1635 1640 1645
 Lys Phe His Leu Lys Arg Ile Arg Glu Gln Phe Pro Gly Thr Glu Met
 1650 1655 1660
 Glu Lys Tyr Ala Leu Phe Thr Tyr Glu Ser Leu Lys Lys Thr Lys Cys
 1665 1670 1675 1680
 Arg Glu Phe Val Pro Ser Arg Asp Glu Ile Glu Ala Leu Ile His Arg
 1685 1690 1695
 Gln Glu Met Thr Ser Thr Val Tyr Cys His Gly Gly Gly Ser Cys Lys
 1700 1705 1710
 Ile Thr Ile Asn Ser His Thr Thr Ala Gly Glu Val Val Glu Lys Leu
 1715 1720 1725
 Ile Arg Gly Leu Ala Met Glu Asp Ser Arg Asn Met Phe Ala Leu Phe
 1730 1735 1740
 Glu Tyr Asn Gly His Val Asp Lys Ala Ile Glu Ser Arg Thr Val Val
 1745 1750 1755 1760
 Ala Asp Val Leu Ala Lys Phe Glu Lys Leu Ala Ala Thr Ser Glu Val
 1765 1770 1775
 Gly Asp Leu Pro Trp Lys Phe Tyr Phe Lys Leu Tyr Cys Phe Leu Asp
 1780 1785 1790
 Thr Asp Asn Val Pro Lys Asp Ser Val Glu Phe Ala Phe Met Phe Glu
 1795 1800 1805
 Gln Ala His Glu Ala Val Ile His Gly His His Pro Ala Pro Glu Glu
 1810 1815 1820
 Asn Leu Gln Val Leu Ala Ala Leu Arg Leu Gln Tyr Leu Gln Gly Asp
 1825 1830 1835 1840

Tyr Thr Leu His Ala Ala Ile Pro Pro Leu Glu Glu Val Tyr Ser Leu
 1845 1850 1855
 Gln Arg Leu Lys Ala Arg Ile Ser Gln Ser Thr Lys Thr Phe Thr Pro
 1860 1865 1870
 Cys Glu Arg Leu Glu Lys Arg Arg Thr Ser Phe Leu Glu Gly Thr Leu
 1875 1880 1885
 Arg Arg Ser Phe Arg Thr Gly Ser Val Val Arg Gln Lys Val Glu Glu
 1890 1895 1900
 Glu Gln Met Leu Asp Met Trp Ile Lys Glu Glu Val Ser Ser Ala Arg
 1905 1910 1915 1920
 Ala Ser Ile Ile Asp Lys Trp Arg Lys Phe Gln Gly Met Asn Gln Glu
 1925 1930 1935
 Gln Ala Met Ala Lys Tyr Met Ala Leu Ile Lys Glu Trp Pro Gly Tyr
 1940 1945 1950
 Gly Ser Thr Leu Phe Asp Val Glu Val Arg Thr Gly Cys His Val Leu
 1955 1960 1965
 Gly Trp Ala Gly Cys Trp His Leu Arg Thr Trp Ile Thr Ala Lys Phe
 1970 1975 1980
 Met Trp Arg Glu Asp Lys Met Glu His Phe Ala Leu Ser Thr Ser Phe
 1985 1990 1995 2000
 Phe Arg Ala Pro Lys Ile Val Pro Leu Thr Pro Pro Phe Ser Ser Gln
 2005 2010 2015
 Phe Leu Phe Ser Cys Val Val Asn Ala Ser Val Ile Leu Gly Met Asn
 2020 2025 2030
 Ala Lys Leu Arg Cys His Leu Phe Phe Tyr Pro Ser Leu Gly Lys Leu
 2035 2040 2045

<210> 13

<211> 1288

<212> DNA

<213> Homo sapiens

<400> 13

ccaacttttg cagctccacc caggatgtgg cctcgctcca cccagctgt gcgcctctct 60
 ccacccttag gcgaaggcac tagaatttcc caaattaaga acgaagagga agtttggacc 120
 ttttcggcca ccgctcgctt caatatggct gccccaggg agagacgagg ctaccatgaa 180
 ggagccgagc gcagaccctg agtccgtcac ccatggatcg cagcgcggag ttcaggaaat 240
 ggaaggcgca atgtttgagc aaagcggacc tcagccggaa gggcagtgtt gacgaggatg 300
 tggtagagct tgtgcagttt ctgaacatgc gagatcagtt tttcaccacc agctccttcg 360
 ctggccgcat cctactcctt gaccggggta taaatggttt tgagggttcag aaacaaaact 420
 gttgctggct actggttaca cacaaacttt gtgtaaaaga tgatgtgatt gtactctga 480
 agaaagcaaa tggatgatgcc actttgaaat ttgaaccatt tgttcttcatt gtgcagtgtc 540
 gacaattgca ggatgcacag attctgcatt ccatggcaat agattctggg ttcaggaaact 600
 ctggcataac ggtgggaaag agaggaaaaa ctatgttggc tgtccggagt acacatggct 660
 tagaagttcc attaagccat aagggaacac tgatggtgac agaggaatat attgacttcc 720

```

tggttaaagt ggcaaataa aaaatggagg aaaacaagaa aagaattgag aggtttttaca 780
actgcctaca gcatgctttg gaaagggaaa cgatgactaa cttacatccc aagatcaaag 840
agaaaaataa ctcatcatat attcataaga aaaaaagaaa cccagaaaaa acacgtgccc 900
agtgtattac taaagaaagt gatgaagaac ttgaaaatga tgatgatgat gatctaggaa 960
tcaatgttac catcttcctt gaagattact aagctttggg tctgatgtgt cttggccgta 1020
atgtttctag taggttttat aaagctgctc ttcataagag tatttttagt tgttgagtgt 1080
atcagccatt cataagccag taatgacaag tgcagagctt caaactataa ctttggttgc 1140
cagaggatgt gcagttgtca tctaagctct cagcagtacc cggcttatcc tacgacttca 1200
cctgaaatgc tatagttatc cctacttttt taccagtttc tcccagaagc acctgcttaa 1260
taaatacaag atgtttgaaa aaaaaaaaa 1288

```

<210> 14

<211> 259

<212> PRT

<213> Homo sapiens

<400> 14

```

Met Asp Arg Ser Ala Glu Phe Arg Lys Trp Lys Ala Gln Cys Leu Ser
  1                      5                      10                      15

Lys Ala Asp Leu Ser Arg Lys Gly Ser Val Asp Glu Asp Val Val Glu
          20                      25                      30

Leu Val Gln Phe Leu Asn Met Arg Asp Gln Phe Phe Thr Thr Ser Ser
          35                      40                      45

Phe Ala Gly Arg Ile Leu Leu Leu Asp Arg Gly Ile Asn Gly Phe Glu
          50                      55                      60

Val Gln Lys Gln Asn Cys Cys Trp Leu Leu Val Thr His Lys Leu Cys
          65                      70                      75

Val Lys Asp Asp Val Ile Val Ala Leu Lys Lys Ala Asn Gly Asp Ala
          85                      90                      95

Thr Leu Lys Phe Glu Pro Phe Val Leu His Val Gln Cys Arg Gln Leu
          100                     105                     110

Gln Asp Ala Gln Ile Leu His Ser Met Ala Ile Asp Ser Gly Phe Arg
          115                     120                     125

Asn Ser Gly Ile Thr Val Gly Lys Arg Gly Lys Thr Met Leu Ala Val
          130                     135                     140

Arg Ser Thr His Gly Leu Glu Val Pro Leu Ser His Lys Gly Lys Leu
          145                     150                     155                     160

Met Val Thr Glu Glu Tyr Ile Asp Phe Leu Leu Asn Val Ala Asn Gln
          165                     170                     175

Lys Met Glu Glu Asn Lys Lys Arg Ile Glu Arg Phe Tyr Asn Cys Leu
          180                     185                     190

Gln His Ala Leu Glu Arg Glu Thr Met Thr Asn Leu His Pro Lys Ile
          195                     200                     205

```

Lys Glu Lys Asn Asn Ser Ser Tyr Ile His Lys Lys Lys Arg Asn Pro
 210 215 220
 Glu Lys Thr Arg Ala Gln Cys Ile Thr Lys Glu Ser Asp Glu Glu Leu
 225 230 235 240
 Glu Asn Asp Asp Asp Asp Asp Leu Gly Ile Asn Val Thr Ile Phe Pro
 245 250 255
 Glu Asp Tyr

<210> 15
 <211> 2352
 <212> DNA
 <213> Homo sapiens
 <220>
 <221> modified_base
 <222> (699)
 <223> a, t, c, g, other or unknown

<400> 15
 acgcgtgcag gtggcgtggc gccagggatt tgaaccgcgc tgacgaagtt tggatgatcca 60
 tcttccgagt atcgccggga tttcgaatcg cgatgatcat cccctctcta gaggagctgg 120
 actccctcaa gtacagtgc ctgcagaact tagccaagag tctgggtctc cgggccaacc 180
 tgagggcaac caagtgttta aaagccttga aaggctacat taaacatgag gcaagaaaaag 240
 gaaatgagaa tcaggatgaa agtcaaactt ctgcatcctc ttgtgatgag actgagatac 300
 agatcagcaa ccaggaagag ctgagagaca gccacttggc catgtcacca aaacaaggag 360
 aaggtgcaag actgtccgtg tggaccctga ctcacagaga atcattcaga gataaaaaata 420
 agtaatccca ctgaattcca gaatcatgaa aagcaggaaa gccaggatct cagagcactg 480
 caaaagtccc ttctccacca gacgagcacc aagaagctga gaatgctgtt tcctcaggta 540
 acagagattc aaaggtacct tcagaaggaa agaaatctct ctacacagat gactcatcca 600
 aacctggaaa aaataaaaaga actgcaatca ctactccaaa ctttaagaag cttcatgaag 660
 ctcatcttaa ggaaatggag tccattgatc caatatatng aggagaaaaa aagaaacatt 720
 ttgaagaaca caattccatg aatgaactga agcagccgcc catcaataag ggaggggtca 780
 ggactccagt acctccaaga ggaagactct ctgtggcttc tactcccatc agccaacgac 840
 gctcgcaagg ccggtcttgt ggccctgcaa gtcagagtac cttgggtctg aaggggtcac 900
 tcaagcgctc tgctatctct gcagctaaaa cgggtgtcag gttttcagct gctactaaag 960
 ataattgagca taagcgttca ctgaccaaga ctccagccag aaagtctgca catgtgaccg 1020
 tgtctggggg cacccaaaaa ggcgaggctg tgcttgggac acacaaatta aagaccatca 1080
 cggggaattc tgctgctgtt attaccccat tcaagttgac aactgaggca acgcagactc 1140
 cagtctccaa taagaaacca gtgtttgatc ttaaagcaag tttgtctcgt cccctcaact 1200
 atgaaccaca caaaggaaag ctaaaaccat gggggcaatc taaagaaaat aattatctaa 1260
 atcaacatgt caacaaatta acttctacaa gaaaacttac aaacaacccc atctccagac 1320
 aaaggaagag caacggaaga aacgcgagca agaagaaagg agaagaaagc aaagggtttg 1380
 ggaatgcgaa ggggcctcat tttggctgaa gattaataat tttttaacat cttgtaataa 1440
 ttctgtatt ctcaactttt ttctttttgt aaattttttt tttttgctgt catccccact 1500
 ttagtcacga gatctttttc tgctaactgt tcatagtctg tgtagtgtcc atgggttctt 1560
 catgtgctat gatctctgaa aagacgttat caccttaaag ctcaaattct ttgggatggg 1620
 ttttacttaa gtccattaac aattcagggt tctaacgaga cccatcctaa aattctcttt 1680
 ctagtttttt aatgtcacca tcccaaacct ccggtttctg atttttaatc cccagctccc 1740
 cagttccctc ttatcgtagt aatattaaca gaactgcagt cttctgctag ccaatagcat 1800
 ttacctgatg gcagctagtt atgcaagctt caggagaatt tgaacaataa caagaatagg 1860
 gtaagctggg atagaaaggc cacctcttca ctctctatag aatatagtaa cctttatgaa 1920
 acggggccat atagtttggt tatgacatca atattttacc taggtgaaat tgtttaggct 1980
 tatgtacctt cgttcaaaata tcctcatgta attgccatct gtcactcact atattcacia 2040


```

aaataaaaact ctacaactca ttctaacatt gcttacttaa aagctacata gccctatcga 2100
aatgcgagga ttaatgcttt aatgctttta gagacagggt ctactgtgt tgcccaggct 2160
ggtctcaaac tccaccaa atgtacttctta ttcattttat ggaaaagact aggctttgct 2220
tagtatcatg tccatgtttc cttcacctca gtggagcttc tgagttttat actgctcaag 2280
atcgtcataa ataaaatttt ttctcattgt caaaaaaaaa aaaaaaaaaa aaaaaaaaaa 2340
aaaaaaaaaa aa 2352

```

<210> 16

<211> 442

<212> PRT

<213> Homo sapiens

<220>

<221> MOD_RES

<222> (204)

<223> Any, other or unknown amino acid

<400> 16

```

Met Ile Ile Pro Ser Leu Glu Glu Leu Asp Ser Leu Lys Tyr Ser Asp
  1             5             10             15

Leu Gln Asn Leu Ala Lys Ser Leu Gly Leu Arg Ala Asn Leu Arg Ala
      20             25             30

Thr Lys Leu Leu Lys Ala Leu Lys Gly Tyr Ile Lys His Glu Ala Arg
      35             40             45

Lys Gly Asn Glu Asn Gln Asp Glu Ser Gln Thr Ser Ala Ser Ser Cys
      50             55             60

Asp Glu Thr Glu Ile Gln Ile Ser Asn Gln Glu Glu Ala Glu Arg Gln
      65             70             75             80

Pro Leu Gly His Val Thr Lys Thr Arg Arg Arg Cys Lys Thr Val Arg
      85             90             95

Val Asp Pro Asp Ser Gln Gln Asn His Ser Glu Ile Lys Ile Ser Asn
      100            105            110

Pro Thr Glu Phe Gln Asn His Glu Lys Gln Glu Ser Gln Asp Leu Arg
      115            120            125

Ala Thr Ala Lys Val Pro Ser Pro Pro Asp Glu His Gln Glu Ala Glu
      130            135            140

Asn Ala Val Ser Ser Gly Asn Arg Asp Ser Lys Val Pro Ser Glu Gly
      145            150            155            160

Lys Lys Ser Leu Tyr Thr Asp Glu Ser Ser Lys Pro Gly Lys Asn Lys
      165            170            175

Arg Thr Ala Ile Thr Thr Pro Asn Phe Lys Lys Leu His Glu Ala His
      180            185            190

Phe Lys Glu Met Glu Ser Ile Asp Pro Ile Tyr Xaa Gly Glu Lys Lys
      195            200            205

```

Lys His Phe Glu Glu His Asn Ser Met Asn Glu Leu Lys Gln Pro Pro
 210 215 220
 Ile Asn Lys Gly Gly Val Arg Thr Pro Val Pro Pro Arg Gly Arg Leu
 225 230 235 240
 Ser Val Ala Ser Thr Pro Ile Ser Gln Arg Arg Ser Gln Gly Arg Ser
 245 250 255
 Cys Gly Pro Ala Ser Gln Ser Thr Leu Gly Leu Lys Gly Ser Leu Lys
 260 265 270
 Arg Ser Ala Ile Ser Ala Ala Lys Thr Gly Val Arg Phe Ser Ala Ala
 275 280 285
 Thr Lys Asp Asn Glu His Lys Arg Ser Leu Thr Lys Thr Pro Ala Arg
 290 295 300
 Lys Ser Ala His Val Thr Val Ser Gly Gly Thr Gln Lys Gly Glu Ala
 305 310 315 320
 Val Leu Gly Thr His Lys Leu Lys Thr Ile Thr Gly Asn Ser Ala Ala
 325 330 335
 Val Ile Thr Pro Phe Lys Leu Thr Thr Glu Ala Thr Gln Thr Pro Val
 340 345 350
 Ser Asn Lys Lys Pro Val Phe Asp Leu Lys Ala Ser Leu Ser Arg Pro
 355 360 365
 Leu Asn Tyr Glu Pro His Lys Gly Lys Leu Lys Pro Trp Gly Gln Ser
 370 375 380
 Lys Glu Asn Asn Tyr Leu Asn Gln His Val Asn Arg Ile Asn Phe Tyr
 385 390 395 400
 Lys Lys Thr Tyr Lys Gln Pro His Leu Gln Thr Lys Glu Glu Gln Arg
 405 410 415
 Lys Lys Arg Glu Gln Glu Arg Lys Glu Lys Lys Ala Lys Val Leu Gly
 420 425 430
 Met Arg Arg Gly Leu Ile Leu Ala Glu Asp
 435 440

<210> 17
 <211> 2058
 <212> PRT
 <213> Homo sapiens

<400> 17
 Met Asp Asn Phe Phe Thr Glu Gly Thr Arg Val Trp Leu Arg Glu Asn
 1 5 10 15
 Gly Gln His Phe Pro Ser Thr Val Asn Ser Cys Ala Glu Gly Ile Val
 20 25 30

Val Phe Arg Thr Asp Tyr Gly Gln Val Phe Thr Tyr Lys Gln Ser Thr
 35 40 45
 Ile Thr His Gln Lys Val Thr Ala Met His Pro Thr Asn Glu Glu Gly
 50 55 60
 Val Asp Asp Met Ala Ser Leu Thr Glu Leu His Gly Gly Ser Ile Met
 65 70 75 80
 Tyr Asn Leu Phe Gln Arg Tyr Lys Arg Asn Gln Ile Tyr Thr Tyr Ile
 85 90 95
 Gly Ser Ile Leu Ala Ser Val Asn Pro Tyr Gln Pro Ile Ala Gly Leu
 100 105 110
 Tyr Glu Pro Ala Thr Met Glu Gln Tyr Ser Arg Arg His Leu Gly Glu
 115 120 125
 Leu Pro Pro His Ile Phe Ala Ile Ala Asn Glu Cys Tyr Arg Cys Leu
 130 135 140
 Trp Lys Arg Tyr Asp Asn Gln Cys Ile Leu Ile Ser Gly Glu Ser Gly
 145 150 155 160
 Ala Gly Lys Thr Glu Ser Thr Lys Leu Ile Leu Lys Phe Leu Ser Val
 165 170 175
 Ile Ser Gln Gln Ser Leu Glu Leu Ser Leu Lys Glu Lys Thr Ser Cys
 180 185 190
 Val Glu Arg Ala Ile Leu Glu Ser Ser Pro Ile Met Glu Ala Phe Gly
 195 200 205
 Asn Ala Lys Thr Val Tyr Asn Asn Asn Ser Ser Arg Phe Gly Lys Phe
 210 215 220
 Val Gln Leu Asn Ile Cys Gln Lys Gly Asn Ile Gln Gly Gly Arg Ile
 225 230 235 240
 Val Asp Tyr Leu Leu Glu Lys Asn Arg Val Val Arg Gln Asn Pro Gly
 245 250 255
 Glu Arg Asn Tyr His Ile Phe Tyr Ala Leu Leu Ala Gly Leu Glu His
 260 265 270
 Glu Glu Arg Glu Glu Phe Tyr Leu Ser Thr Pro Glu Asn Tyr His Tyr
 275 280 285
 Leu Asn Gln Ser Gly Cys Val Glu Asp Lys Thr Ile Ser Asp Gln Glu
 290 295 300
 Ser Phe Arg Glu Val Ile Thr Ala Met Asp Val Met Gln Phe Ser Lys
 305 310 315 320
 Glu Glu Val Arg Glu Val Ser Arg Leu Leu Ala Gly Ile Leu His Leu
 325 330 335

Gly Asn Ile Glu Phe Ile Thr Ala Gly Gly Ala Gln Val Ser Phe Lys
 340 345 350
 Thr Ala Leu Gly Arg Ser Ala Glu Leu Leu Gly Leu Asp Pro Thr Gln
 355 360 365
 Leu Thr Asp Ala Leu Thr Gln Arg Ser Met Phe Leu Arg Gly Glu Glu
 370 375 380
 Ile Leu Thr Pro Leu Asn Val Gln Gln Ala Val Asp Ser Arg Asp Ser
 385 390 395 400
 Leu Ala Met Ala Leu Tyr Ala Cys Cys Phe Glu Trp Val Ile Lys Lys
 405 410 415
 Ile Asn Ser Arg Ile Lys Gly Asn Glu Asp Phe Lys Ser Ile Gly Ile
 420 425 430
 Leu Asp Ile Phe Gly Phe Glu Asn Phe Glu Val Asn His Phe Glu Gln
 435 440 445
 Phe Asn Ile Asn Tyr Ala Asn Glu Lys Leu Gln Glu Tyr Phe Asn Lys
 450 455 460
 His Ile Phe Ser Leu Glu Gln Leu Glu Tyr Ser Arg Glu Gly Leu Val
 465 470 475 480
 Trp Glu Asp Ile Asp Trp Ile Asp Asn Gly Glu Cys Leu Asp Leu Ile
 485 490 495
 Glu Lys Lys Leu Gly Leu Leu Ala Leu Ile Asn Glu Glu Ser His Phe
 500 505 510
 Pro Gln Ala Thr Asp Ser Thr Leu Leu Glu Lys Leu His Ser Gln His
 515 520 525
 Ala Asn Asn His Phe Tyr Val Lys Pro Arg Val Ala Val Asn Asn Phe
 530 535 540
 Gly Val Lys His Tyr Ala Gly Glu Val Gln Tyr Asp Val Arg Gly Ile
 545 550 555 560
 Leu Glu Lys Asn Arg Asp Thr Phe Arg Asp Asp Leu Leu Asn Leu Leu
 565 570 575
 Arg Glu Ser Arg Phe Asp Phe Ile Tyr Asp Leu Phe Glu His Val Ser
 580 585 590
 Ser Arg Asn Asn Gln Asp Thr Leu Lys Cys Gly Ser Lys His Arg Arg
 595 600 605
 Pro Thr Val Ser Ser Gln Phe Lys Asp Ser Leu His Ser Leu Met Ala
 610 615 620
 Thr Leu Ser Ser Ser Asn Pro Phe Phe Val Arg Cys Ile Lys Pro Asn
 625 630 635 640

Met Gln Lys Met Pro Asp Gln Phe Asp Gln Ala Val Val Leu Asn Gln
 645 650 655
 Leu Arg Tyr Ser Gly Met Leu Glu Thr Val Arg Ile Arg Lys Ala Gly
 660 665 670
 Tyr Ala Val Arg Arg Pro Phe Gln Asp Phe Tyr Lys Arg Tyr Lys Val
 675 680 685
 Leu Met Arg Asn Leu Ala Leu Pro Glu Asp Val Arg Gly Lys Cys Thr
 690 695 700
 Ser Leu Leu Gln Leu Tyr Asp Ala Ser Asn Ser Glu Trp Gln Leu Gly
 705 710 715 720
 Lys Thr Lys Val Phe Leu Arg Glu Ser Leu Glu Gln Lys Leu Glu Lys
 725 730 735
 Arg Arg Glu Glu Glu Val Ser His Ala Ala Met Val Ile Arg Ala His
 740 745 750
 Val Leu Gly Phe Leu Ala Arg Lys Gln Tyr Arg Lys Val Leu Tyr Cys
 755 760 765
 Val Val Ile Ile Gln Lys Asn Tyr Arg Ala Phe Leu Leu Arg Arg Arg
 770 775 780
 Phe Leu His Leu Lys Lys Ala Ala Ile Val Phe Gln Lys Gln Leu Arg
 785 790 795 800
 Gly Gln Ile Ala Arg Arg Val Tyr Arg Gln Leu Leu Ala Glu Lys Arg
 805 810 815
 Glu Gln Glu Glu Lys Lys Lys Gln Glu Glu Glu Glu Lys Lys Lys Arg
 820 825 830
 Glu Glu Glu Glu Arg Glu Arg Glu Arg Glu Arg Arg Glu Ala Glu Leu
 835 840 845
 Arg Ala Gln Gln Glu Glu Glu Thr Arg Lys Gln Gln Glu Leu Glu Ala
 850 855 860
 Leu Gln Lys Ser Gln Lys Glu Ala Glu Leu Thr Arg Glu Leu Glu Lys
 865 870 875 880
 Gln Lys Glu Asn Lys Gln Val Glu Glu Ile Leu Arg Leu Glu Lys Glu
 885 890 895
 Ile Glu Asp Leu Gln Arg Met Lys Glu Gln Gln Glu Leu Ser Leu Thr
 900 905 910
 Glu Ala Ser Leu Gln Lys Leu Gln Glu Arg Arg Asp Gln Glu Leu Arg
 915 920 925
 Arg Leu Glu Glu Glu Ala Cys Arg Ala Ala Gln Glu Phe Leu Glu Ser
 930 935 940

Leu Asn Phe Asp Glu Ile Asp Glu Cys Val Arg Asn Ile Glu Arg Ser
 945 950 955 960
 Leu Ser Val Gly Ser Glu Phe Ser Ser Glu Leu Ala Glu Ser Ala Cys
 965 970 975
 Glu Glu Lys Pro Asn Phe Asn Phe Ser Gln Pro Tyr Pro Glu Glu Glu
 980 985 990
 Val Asp Glu Gly Phe Glu Ala Asp Asp Asp Ala Phe Lys Asp Ser Pro
 995 1000 1005
 Asn Pro Ser Glu His Gly His Ser Asp Gln Arg Thr Ser Gly Ile Arg
 1010 1015 1020
 Thr Ser Asp Asp Ser Ser Glu Glu Asp Pro Tyr Met Asn Asp Thr Val
 1025 1030 1035 1040
 Val Pro Thr Ser Pro Ser Ala Asp Ser Thr Val Leu Leu Ala Pro Ser
 1045 1050 1055
 Val Gln Asp Ser Gly Ser Leu His Asn Ser Ser Ser Gly Glu Ser Thr
 1060 1065 1070
 Tyr Cys Met Pro Gln Asn Ala Gly Asp Leu Pro Ser Pro Asp Gly Asp
 1075 1080 1085
 Tyr Asp Tyr Asp Gln Asp Asp Tyr Glu Asp Gly Ala Ile Thr Ser Gly
 1090 1095 1100
 Ser Ser Val Thr Phe Ser Asn Ser Tyr Gly Ser Gln Trp Ser Pro Asp
 1105 1110 1115 1120
 Tyr Arg Cys Ser Val Gly Thr Tyr Asn Ser Ser Gly Ala Tyr Arg Phe
 1125 1130 1135
 Ser Ser Glu Gly Ala Gln Ser Ser Phe Glu Asp Ser Glu Glu Asp Phe
 1140 1145 1150
 Asp Ser Arg Phe Asp Thr Asp Asp Glu Leu Ser Tyr Arg Arg Asp Ser
 1155 1160 1165
 Val Tyr Ser Cys Val Thr Leu Pro Tyr Phe His Ser Phe Leu Tyr Met
 1170 1175 1180
 Lys Gly Gly Leu Met Asn Ser Trp Lys Arg Arg Trp Cys Val Leu Lys
 1185 1190 1195 1200
 Asp Glu Thr Phe Leu Trp Phe Arg Ser Lys Gln Glu Ala Leu Lys Gln
 1205 1210 1215
 Gly Trp Leu His Lys Lys Gly Gly Gly Ser Ser Thr Leu Ser Arg Arg
 1220 1225 1230
 Asn Trp Lys Lys Arg Trp Phe Val Leu Arg Gln Ser Lys Leu Met Tyr
 1235 1240 1245

Phe Glu Asn Asp Ser Glu Glu Lys Leu Lys Gly Thr Val Glu Val Arg
 1250 1255 1260
 Thr Ala Lys Glu Ile Ile Asp Asn Thr Thr Lys Glu Asn Gly Ile Asp
 1265 1270 1275 1280
 Ile Ile Met Ala Asp Arg Thr Phe His Leu Ile Ala Glu Ser Pro Glu
 1285 1290 1295
 Asp Ala Ser Gln Trp Phe Ser Val Leu Ser Gln Val His Ala Ser Thr
 1300 1305 1310
 Asp Gln Glu Ile Gln Glu Met His Asp Glu Gln Ala Asn Pro Gln Asn
 1315 1320 1325
 Ala Val Gly Thr Leu Asp Val Gly Leu Ile Asp Ser Val Cys Ala Ser
 1330 1335 1340
 Asp Ser Pro Asp Arg Pro Asn Ser Phe Val Ile Ile Thr Ala Asn Arg
 1345 1350 1355 1360
 Val Leu His Cys Asn Ala Asp Thr Pro Glu Glu Met His His Trp Ile
 1365 1370 1375
 Thr Leu Leu Gln Arg Ser Lys Gly Asp Thr Arg Val Glu Gly Gln Glu
 1380 1385 1390
 Phe Ile Val Arg Gly Trp Leu His Lys Glu Val Lys Asn Ser Pro Lys
 1395 1400 1405
 Met Ser Ser Leu Lys Leu Lys Lys Arg Trp Phe Val Leu Thr His Asn
 1410 1415 1420
 Ser Leu Asp Tyr Tyr Lys Ser Ser Glu Lys Asn Ala Leu Lys Leu Gly
 1425 1430 1435 1440
 Thr Leu Val Leu Asn Ser Leu Cys Ser Val Val Pro Pro Asp Glu Lys
 1445 1450 1455
 Ile Phe Lys Glu Thr Gly Tyr Trp Asn Val Thr Val Tyr Gly Arg Lys
 1460 1465 1470
 His Cys Tyr Arg Leu Tyr Thr Lys Leu Leu Asn Glu Ala Thr Arg Trp
 1475 1480 1485
 Ser Ser Ala Ile Gln Asn Val Thr Asp Thr Lys Ala Pro Ile Asp Thr
 1490 1495 1500
 Pro Thr Gln Gln Leu Ile Gln Asp Ile Lys Glu Asn Cys Leu Asn Ser
 1505 1510 1515 1520
 Asp Val Val Glu Gln Ile Tyr Lys Arg Asn Pro Ile Leu Arg Tyr Thr
 1525 1530 1535
 His His Pro Leu His Ser Pro Leu Leu Pro Leu Pro Tyr Gly Asp Ile
 1540 1545 1550

Asn Leu Asn Leu Leu Lys Asp Lys Gly Tyr Thr Thr Leu Gln Asp Glu
 1555 1560 1565
 Ala Ile Lys Ile Phe Asn Ser Leu Gln Gln Leu Glu Ser Met Ser Asp
 1570 1575 1580
 Pro Ile Pro Ile Ile Gln Gly Ile Leu Gln Thr Gly His Asp Leu Arg
 1585 1590 1595 1600
 Pro Leu Arg Asp Glu Leu Tyr Cys Gln Leu Ile Lys Gln Thr Asn Lys
 1605 1610 1615
 Val Pro His Pro Gly Ser Val Gly Asn Leu Tyr Ser Trp Gln Ile Leu
 1620 1625 1630
 Thr Cys Leu Ser Cys Thr Phe Leu Pro Ser Arg Gly Ile Leu Lys Tyr
 1635 1640 1645
 Leu Lys Phe His Leu Lys Arg Ile Arg Glu Gln Phe Pro Gly Thr Glu
 1650 1655 1660
 Met Glu Lys Tyr Ala Leu Phe Thr Tyr Glu Ser Leu Lys Lys Thr Lys
 1665 1670 1675 1680
 Cys Arg Glu Phe Val Pro Ser Arg Asp Glu Ile Glu Ala Leu Ile His
 1685 1690 1695
 Arg Gln Glu Met Thr Ser Thr Val Tyr Cys His Gly Gly Gly Ser Cys
 1700 1705 1710
 Lys Ile Thr Ile Asn Ser His Thr Thr Ala Gly Glu Val Val Glu Lys
 1715 1720 1725
 Leu Ile Arg Gly Leu Ala Met Glu Asp Ser Arg Asn Met Phe Ala Leu
 1730 1735 1740
 Phe Glu Tyr Asn Gly His Val Asp Lys Ala Ile Glu Ser Arg Thr Val
 1745 1750 1755 1760
 Val Ala Asp Val Leu Ala Lys Phe Glu Lys Leu Ala Ala Thr Ser Glu
 1765 1770 1775
 Val Gly Asp Leu Pro Trp Lys Phe Tyr Phe Lys Leu Tyr Cys Phe Leu
 1780 1785 1790
 Asp Thr Asp Asn Val Pro Lys Asp Ser Val Glu Phe Ala Phe Met Phe
 1795 1800 1805
 Glu Gln Ala His Glu Ala Val Ile His Gly His His Pro Ala Pro Glu
 1810 1815 1820
 Glu Asn Leu Gln Val Leu Ala Ala Leu Arg Leu Gln Tyr Leu Gln Gly
 1825 1830 1835 1840
 Asp Tyr Thr Leu His Ala Ala Ile Pro Pro Leu Glu Glu Val Tyr Ser
 1845 1850 1855

Leu Gln Arg Leu Lys Ala Arg Ile Ser Gln Ser Thr Lys Thr Phe Thr
 1860 1865 1870
 Pro Cys Glu Arg Leu Glu Lys Arg Arg Thr Ser Phe Leu Glu Gly Thr
 1875 1880 1885
 Leu Arg Arg Ser Phe Arg Thr Gly Ser Val Val Arg Gln Lys Val Glu
 1890 1895 1900
 Glu Glu Gln Met Leu Asp Met Trp Ile Lys Glu Glu Val Ser Ser Ala
 1905 1910 1915 1920
 Arg Ala Ser Ile Ile Asp Lys Trp Arg Lys Phe Gln Gly Met Asn Gln
 1925 1930 1935
 Glu Gln Ala Met Ala Lys Tyr Met Ala Leu Ile Lys Glu Trp Pro Gly
 1940 1945 1950
 Tyr Gly Ser Thr Leu Phe Asp Val Glu Cys Lys Glu Gly Gly Phe Pro
 1955 1960 1965
 Gln Glu Leu Trp Leu Gly Val Ser Ala Asp Ala Val Ser Val Tyr Lys
 1970 1975 1980
 Arg Gly Glu Gly Arg Pro Leu Glu Val Phe Gln Tyr Glu His Ile Leu
 1985 1990 1995 2000
 Ser Phe Gly Ala Pro Leu Ala Asn Thr Tyr Lys Ile Val Val Asp Glu
 2005 2010 2015
 Arg Glu Leu Leu Phe Glu Thr Ser Glu Val Val Asp Val Ala Lys Leu
 2020 2025 2030
 Met Lys Ala Tyr Ile Ser Met Ile Val Lys Lys Arg Tyr Ser Thr Thr
 2035 2040 2045
 Arg Ser Ala Ser Ser Gln Gly Ser Ser Arg
 2050 2055